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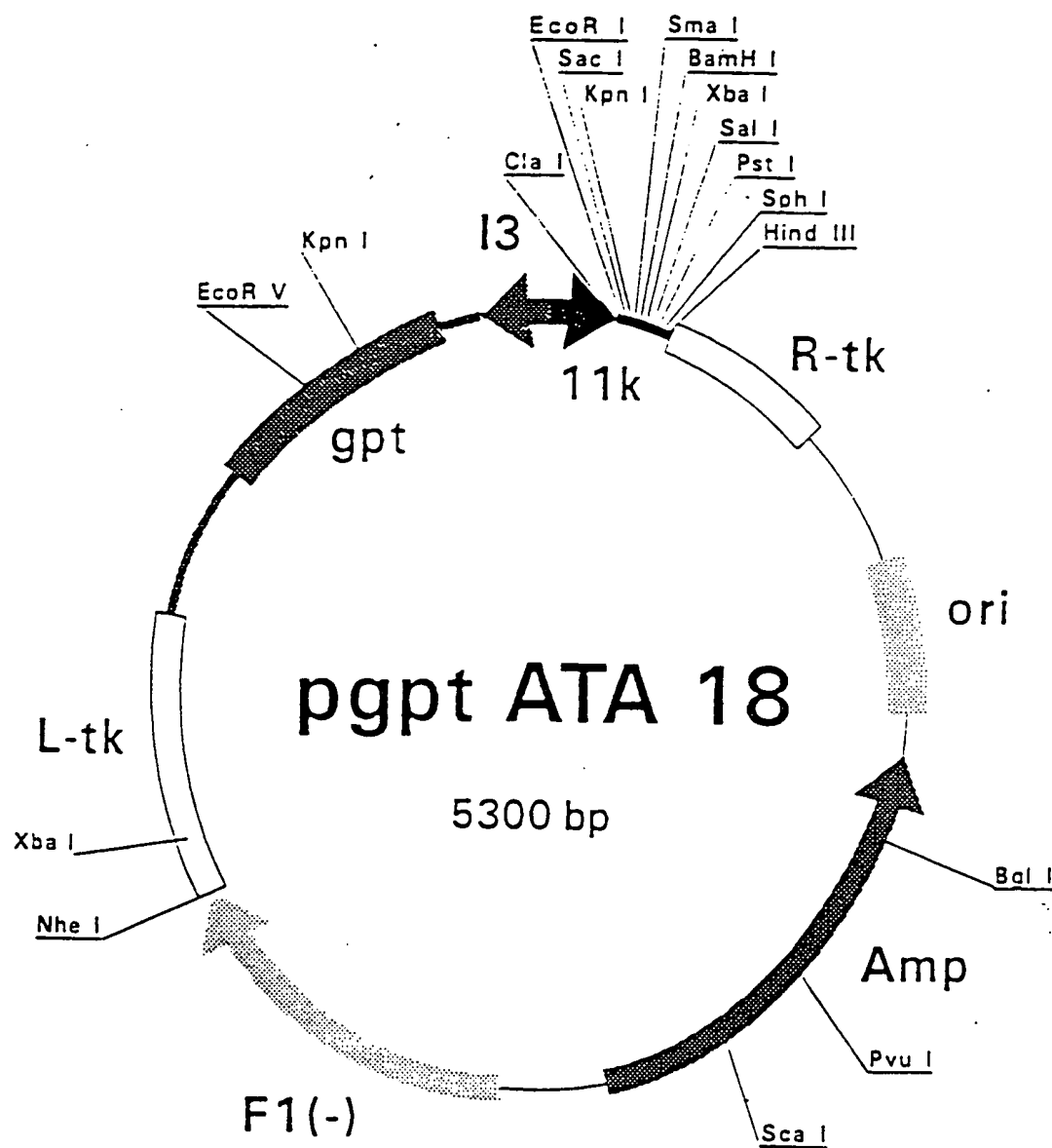


FIGURE 1

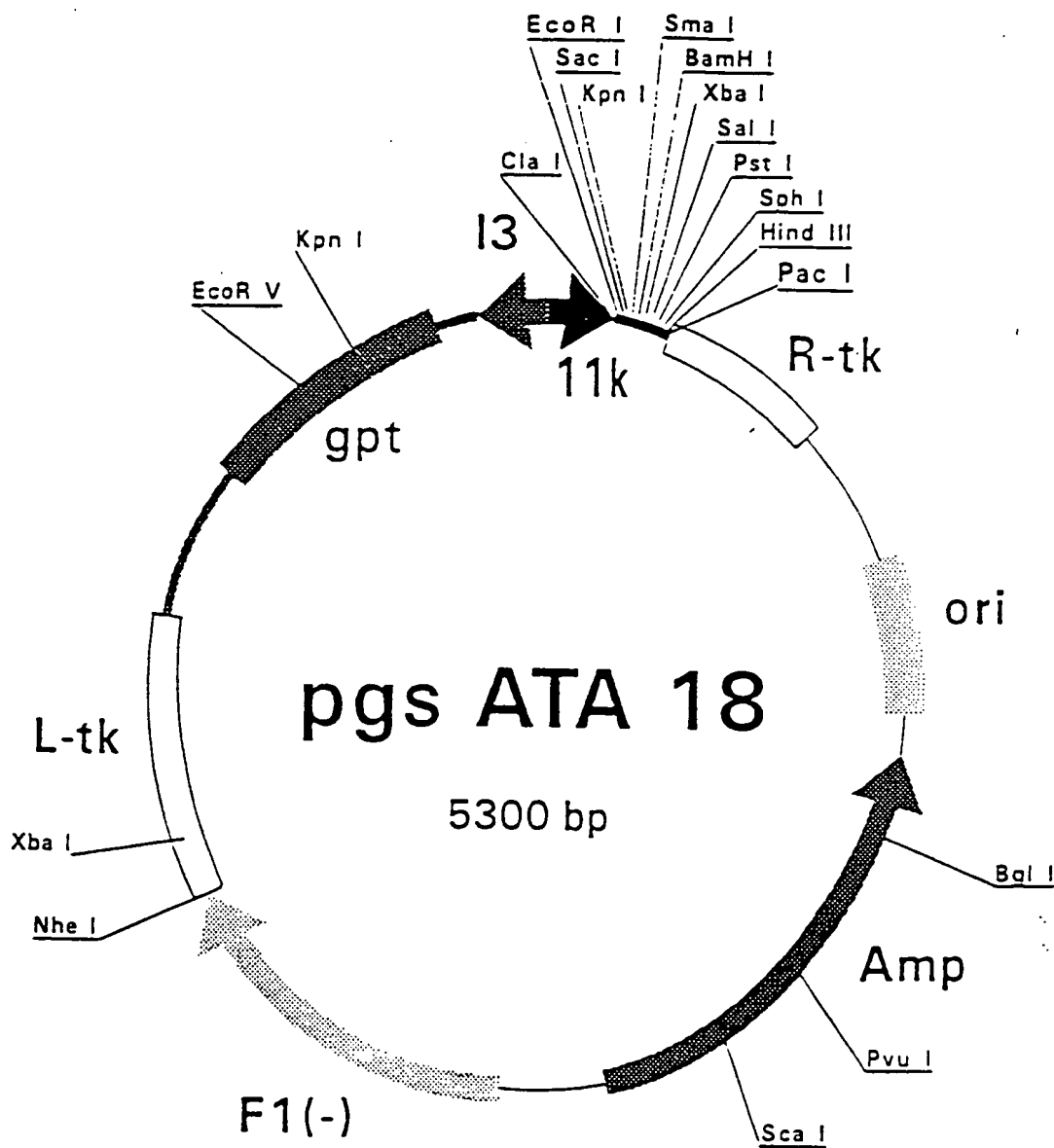


FIGURE 2

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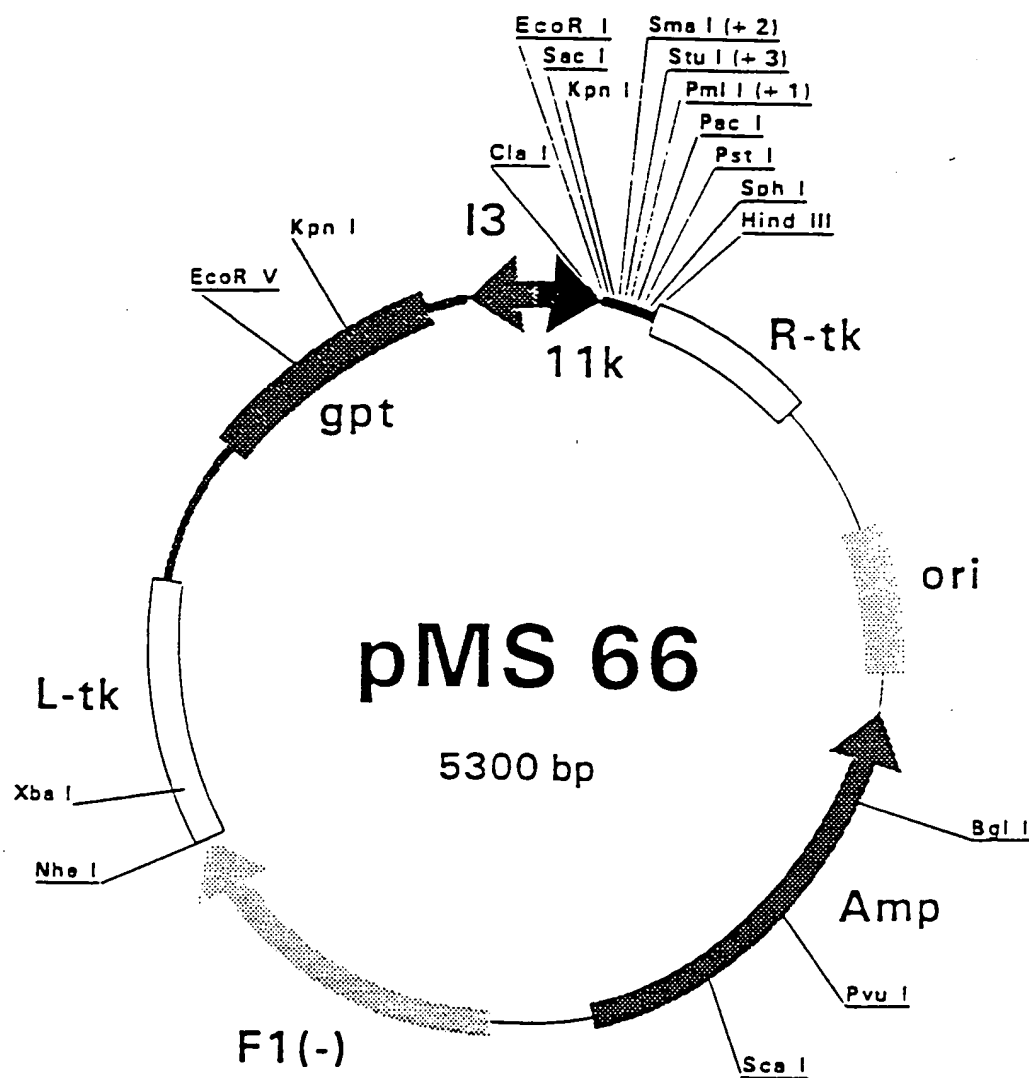


FIGURE 3

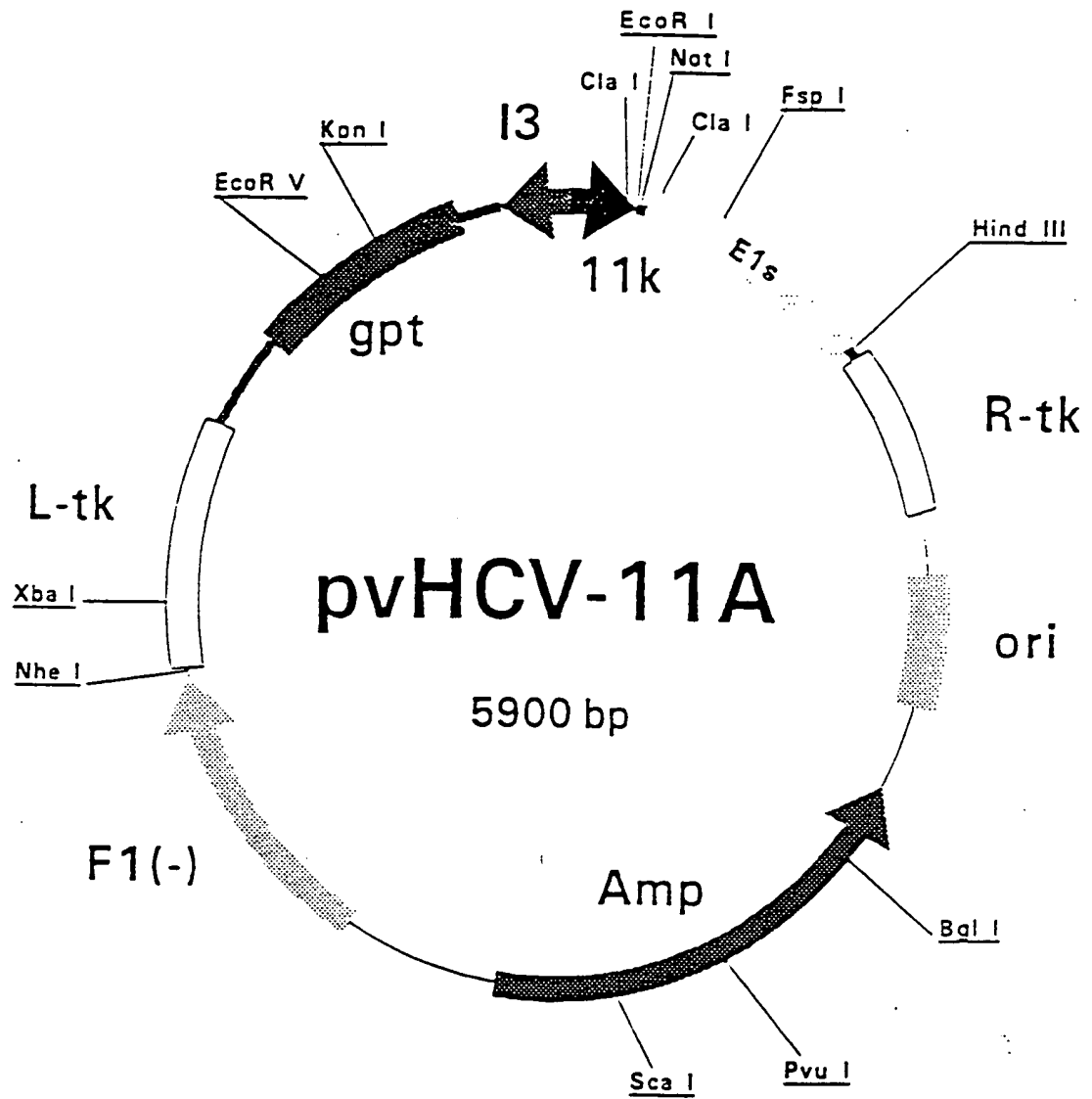


FIGURE 4

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Anti-E1 levels in NON-responders to IFN treatment

Series 1

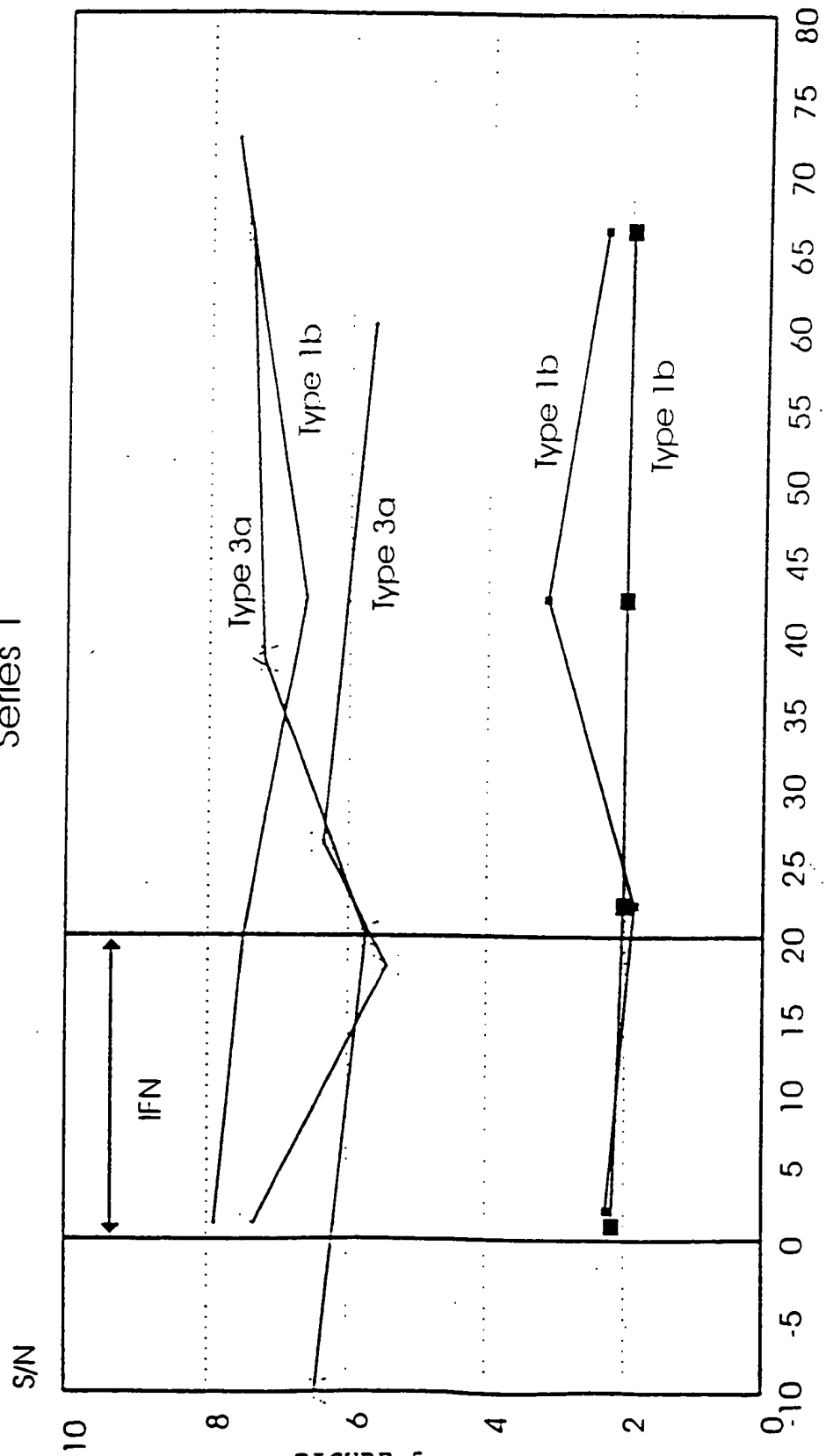
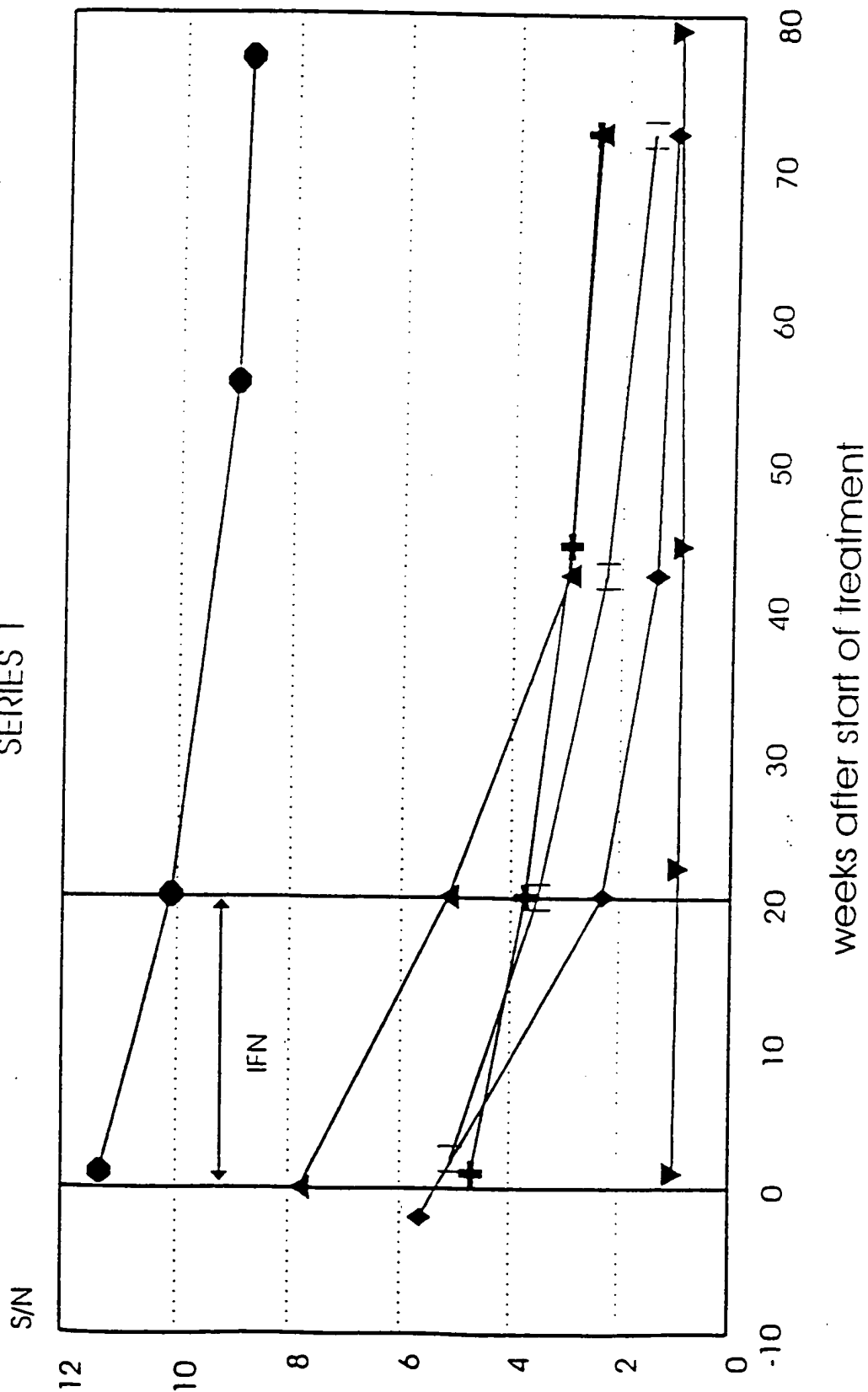


FIGURE 5

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Anti-E1 levels in RESPONDERS to IFN treatment

SERIES 1



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Anti-E1 levels in patients with COMPLETE response to IFN

SERIES 2

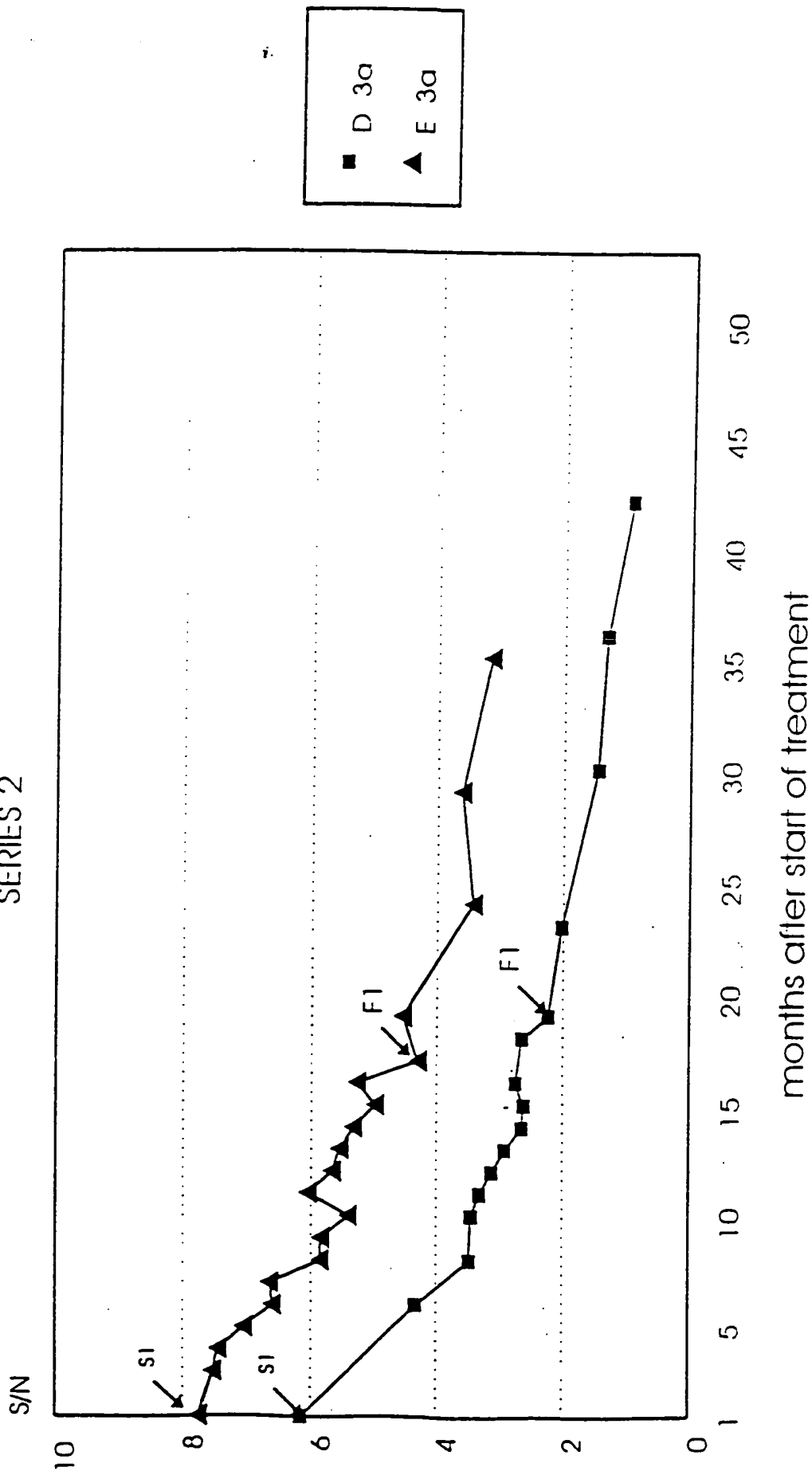


FIGURE 7

Anti-E1 levels in INCOMPLETE responders to IFN treatment

SERIES 2

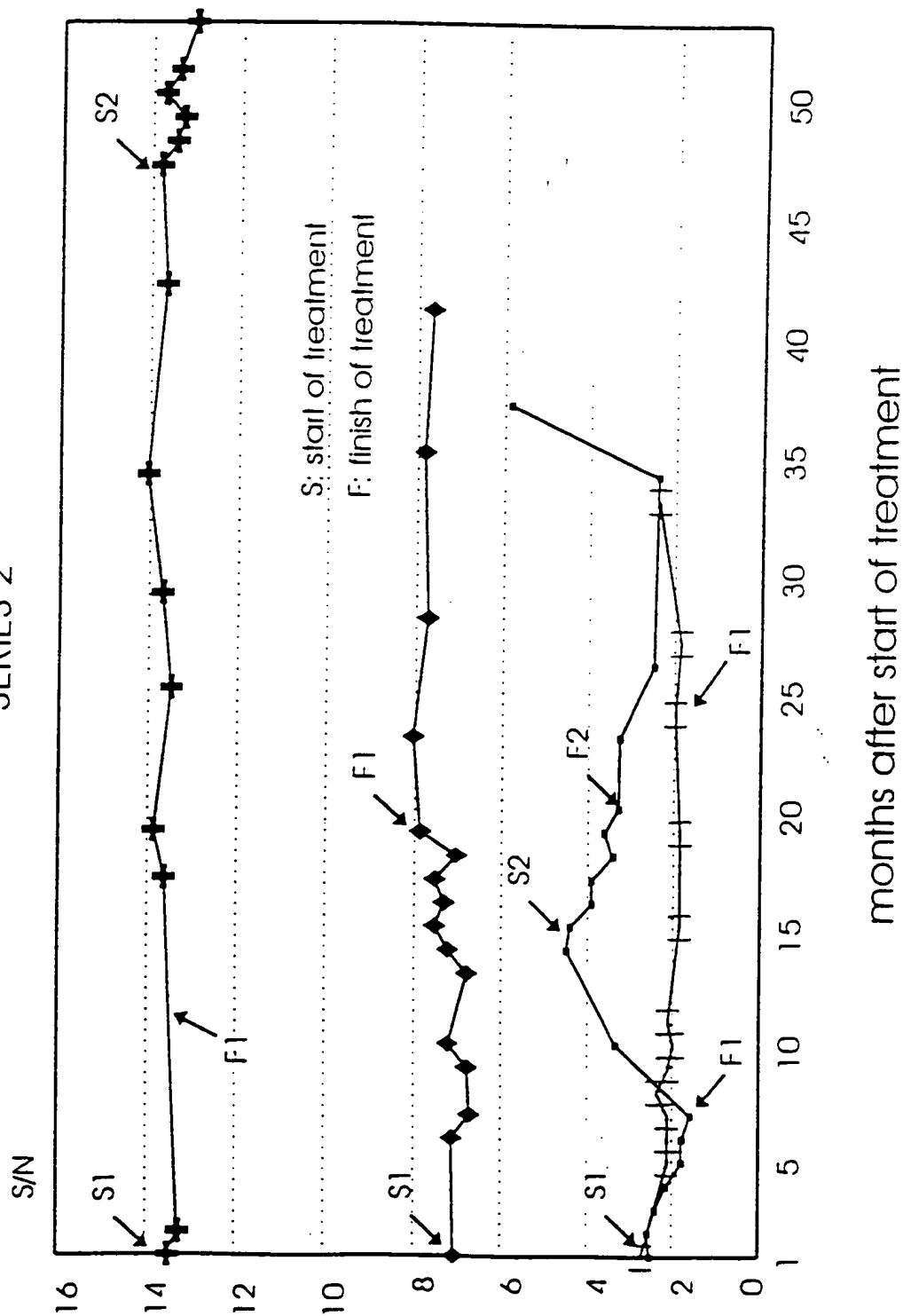


FIGURE 2

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Anti-E2 levels in NON-RESPONDERS to IFN treatment

SERIES 1

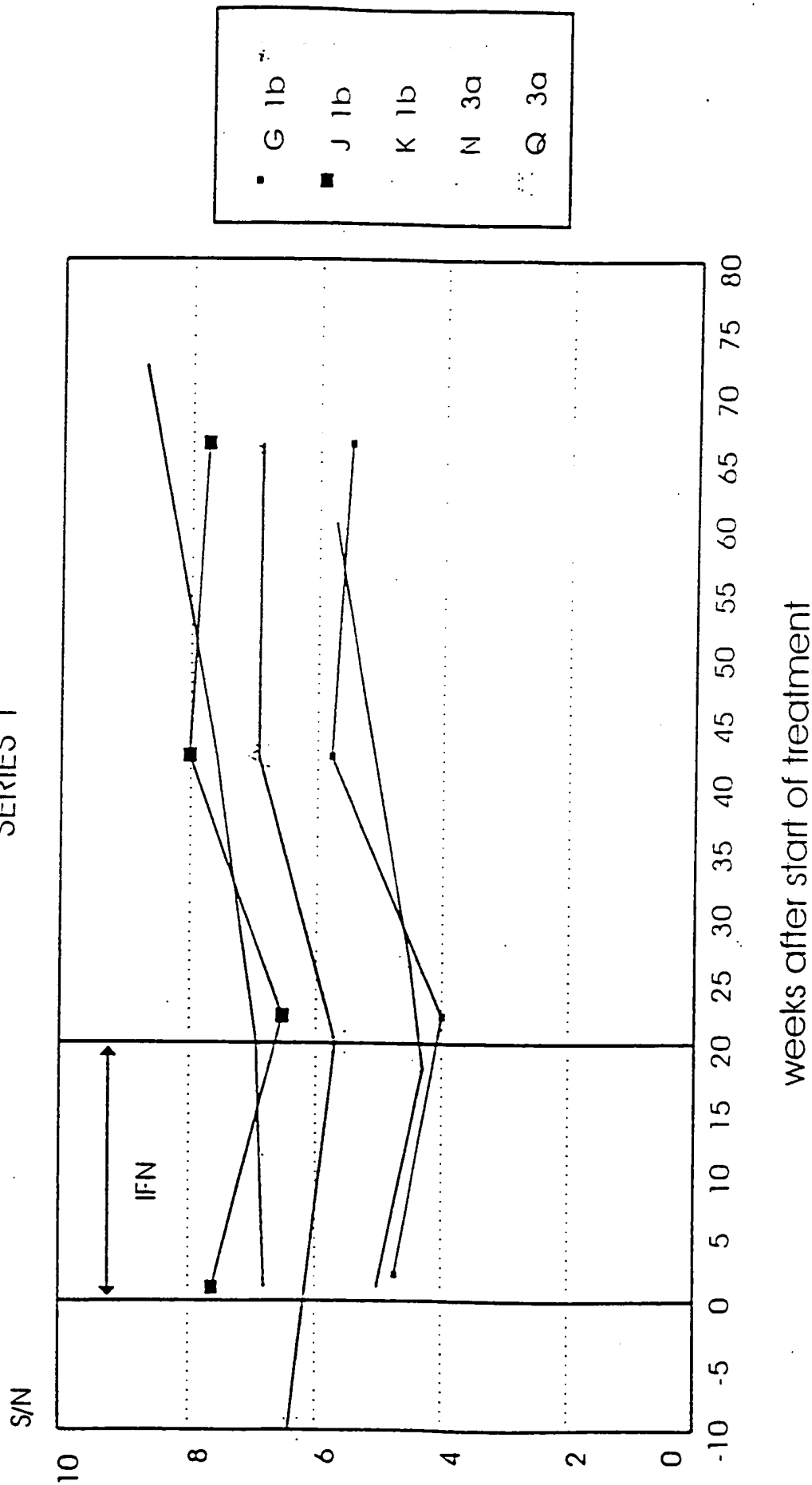
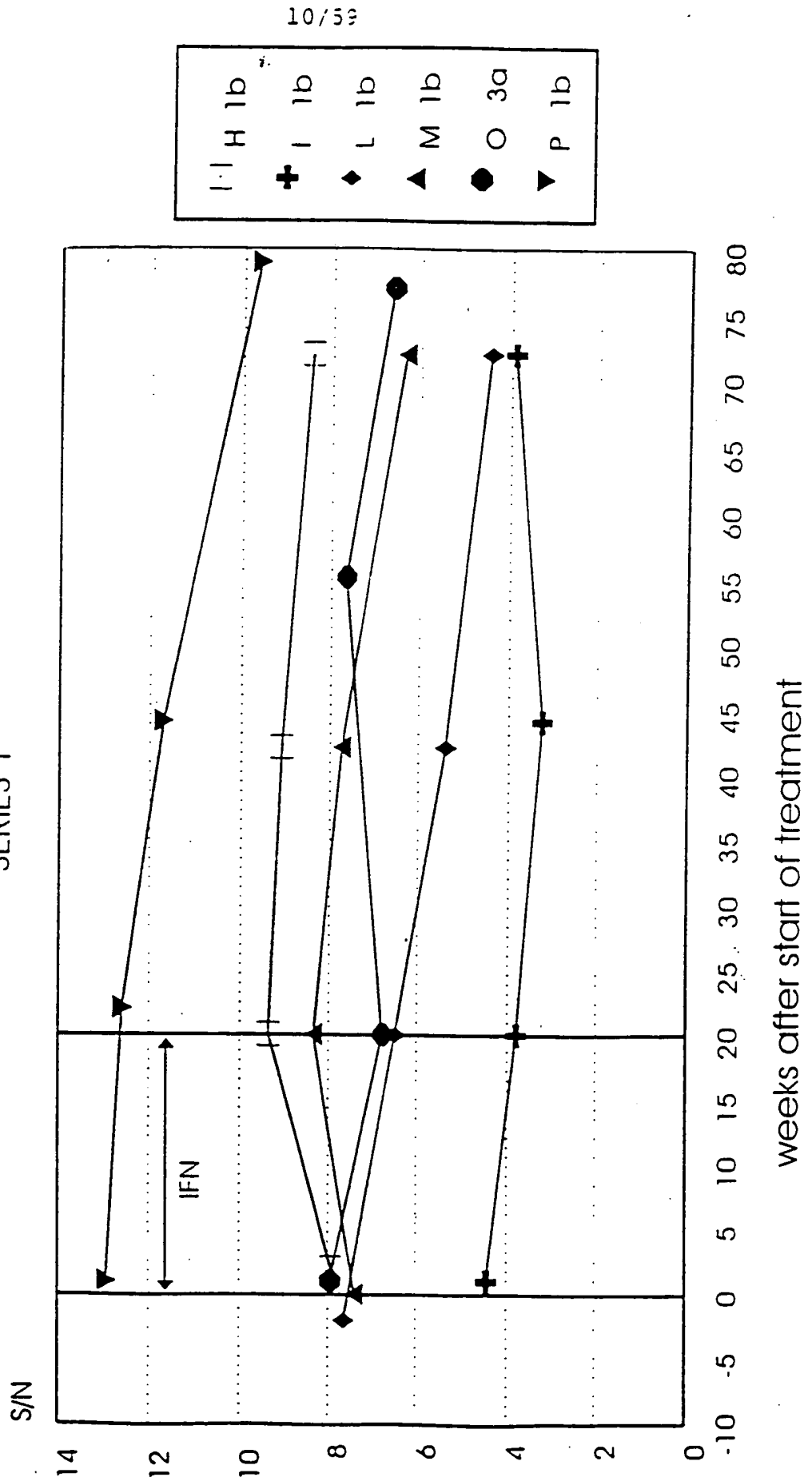


FIGURE 9

Anti-E2 levels in RESPONDERS to IFN treatment

SERIES 1



Anti-E2 levels in INCOMPLETE responders to IFN treatment

SERIES 2

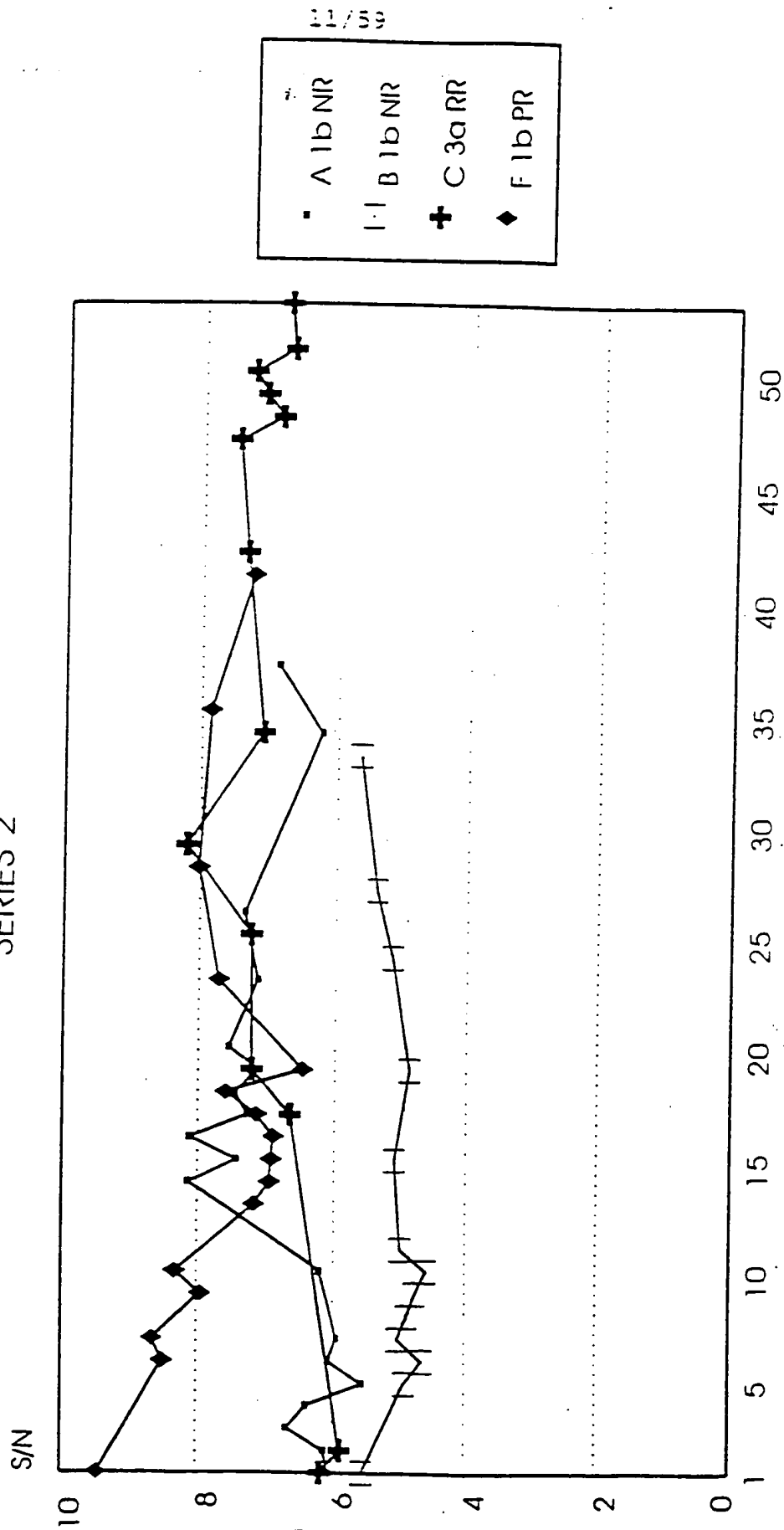


FIGURE 11

months after start of treatment

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Anti-E2 levels in COMPLETE responders to IFN treatment

SERIES 2

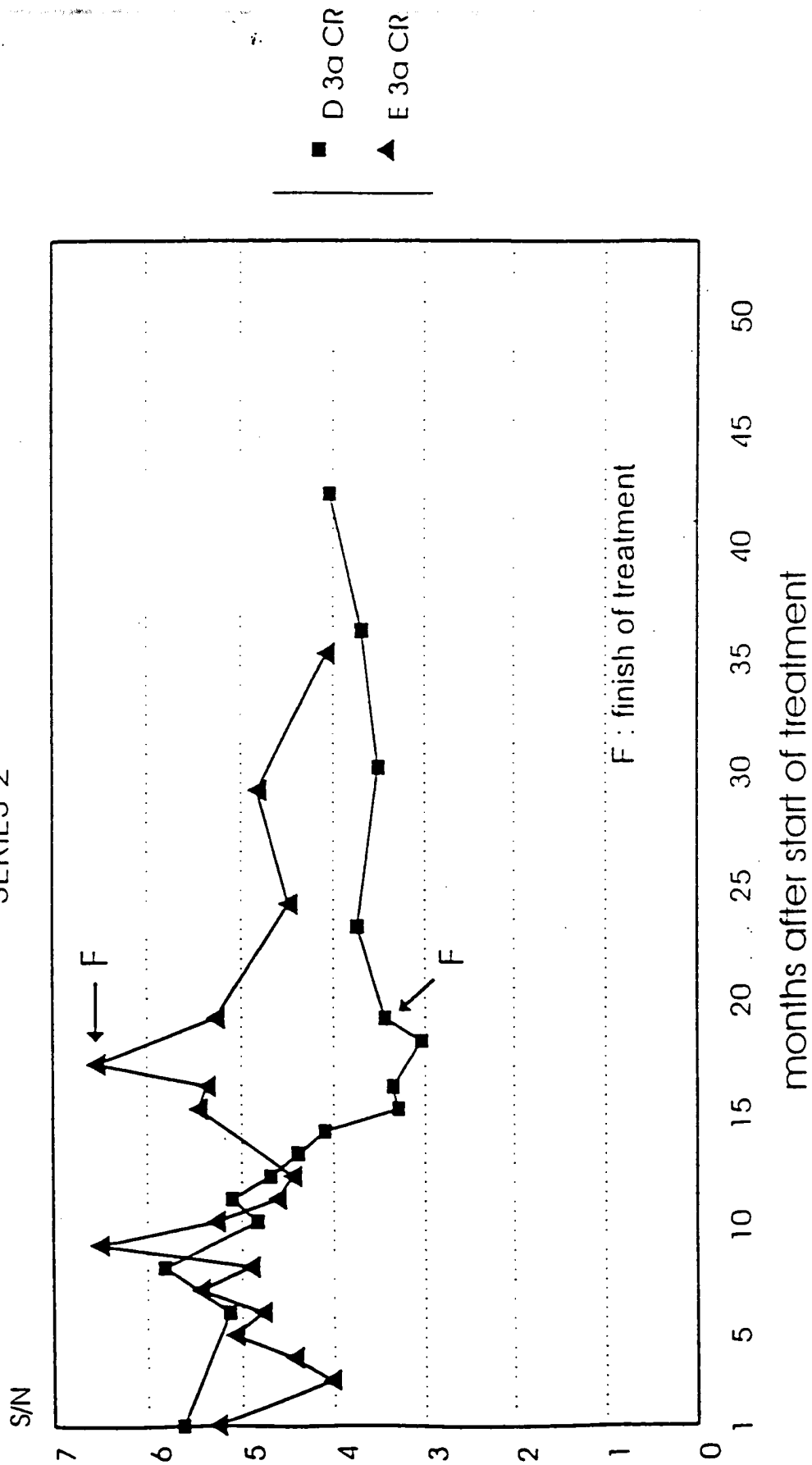


FIGURE 1'

FIGURE 13

Human anti-E1 reactivity competed with peptides

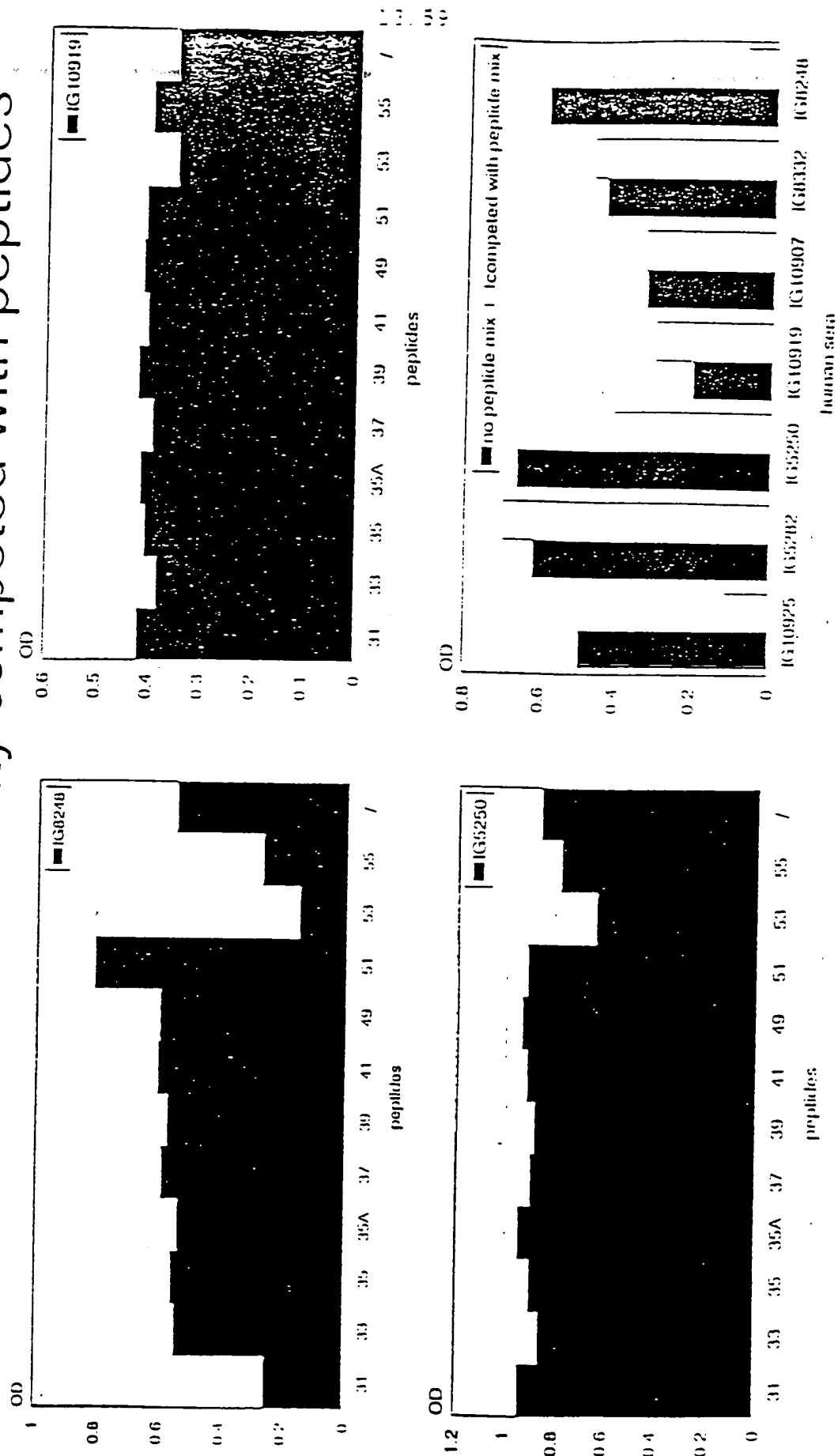
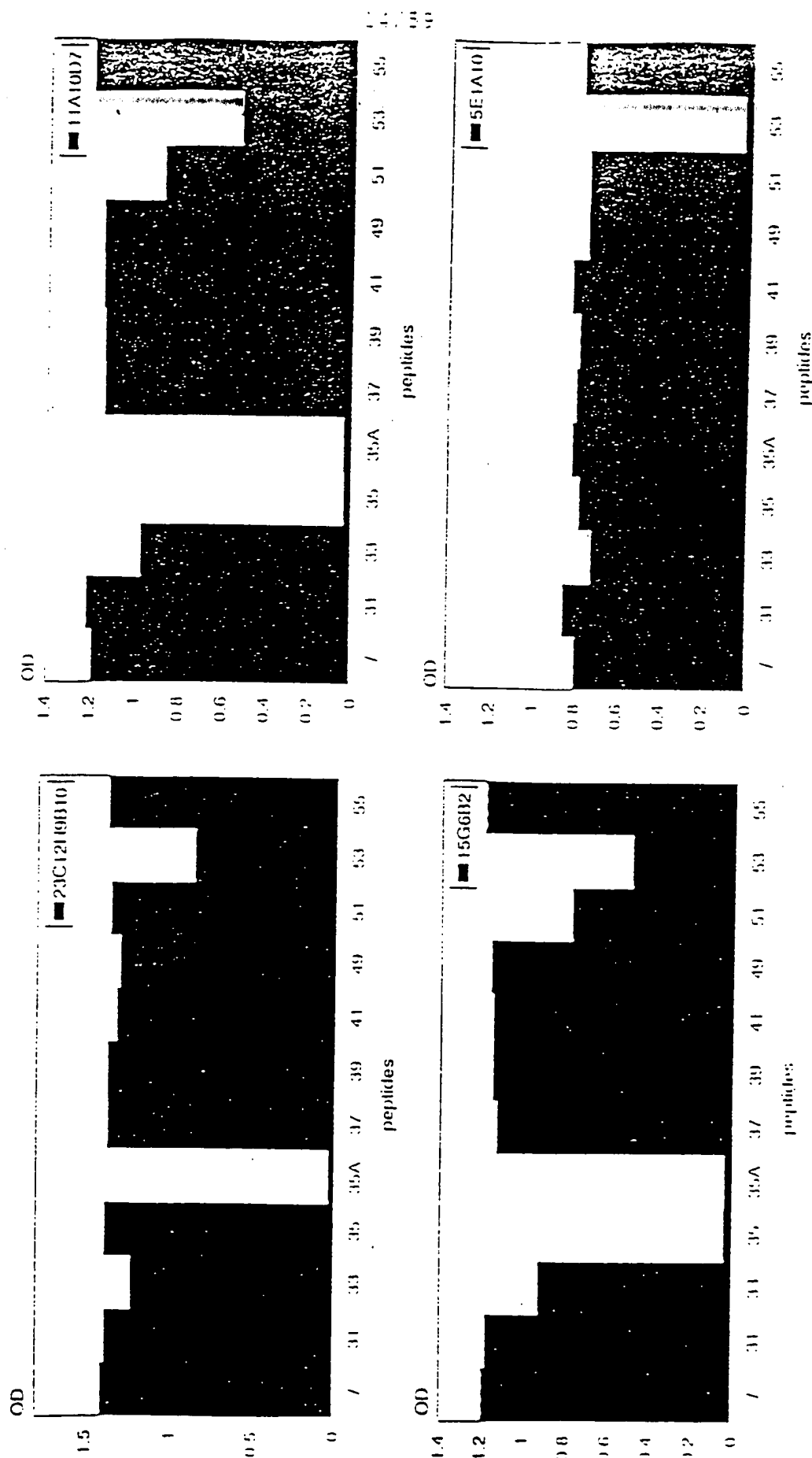


FIGURE 14

Competition of reactivity of anti-E1 Mabs with peptides



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Anti-E1 (epitope 1) levels in NON-RESPONDERS to IFN treatment

SERIES 1

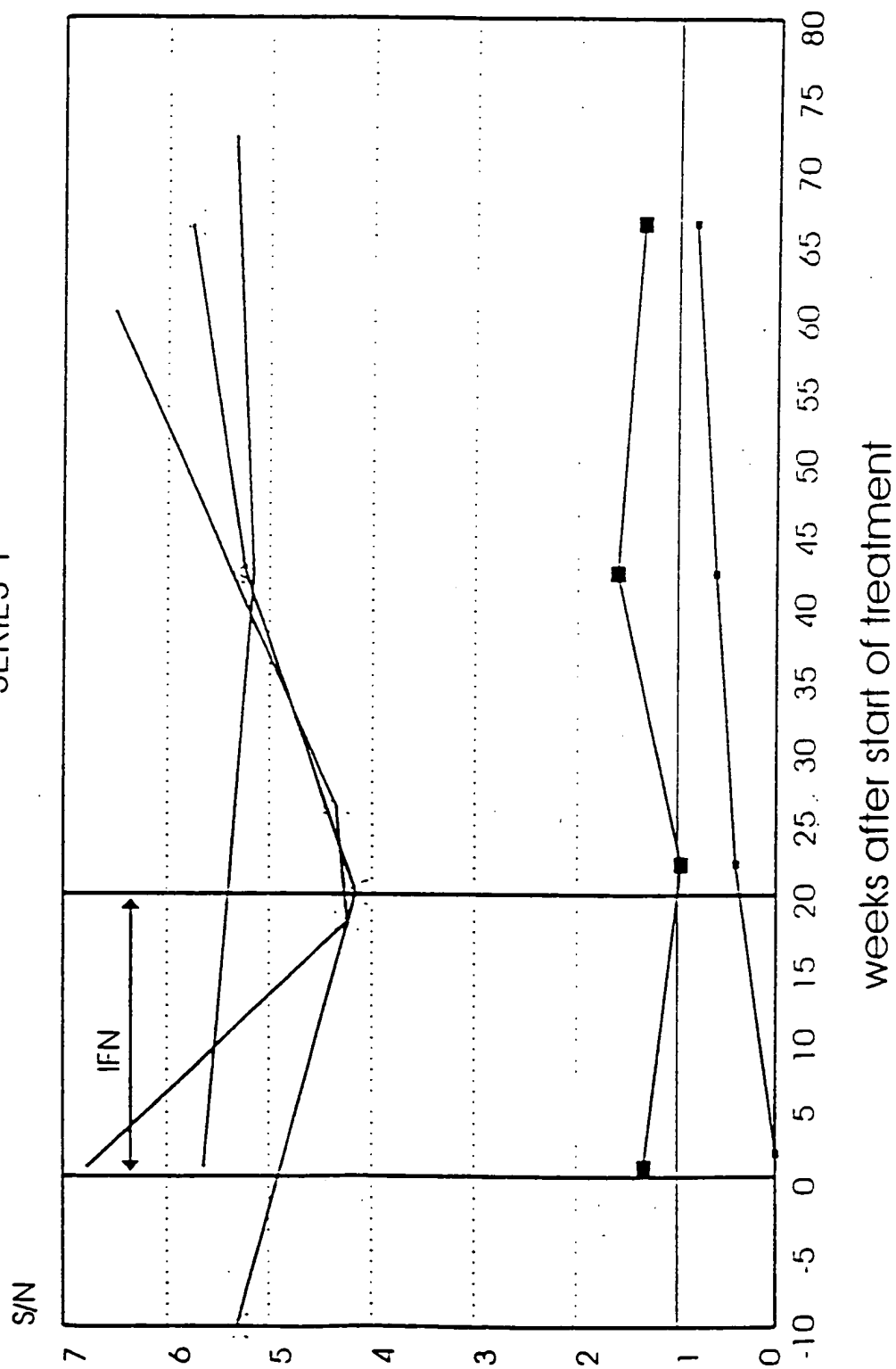
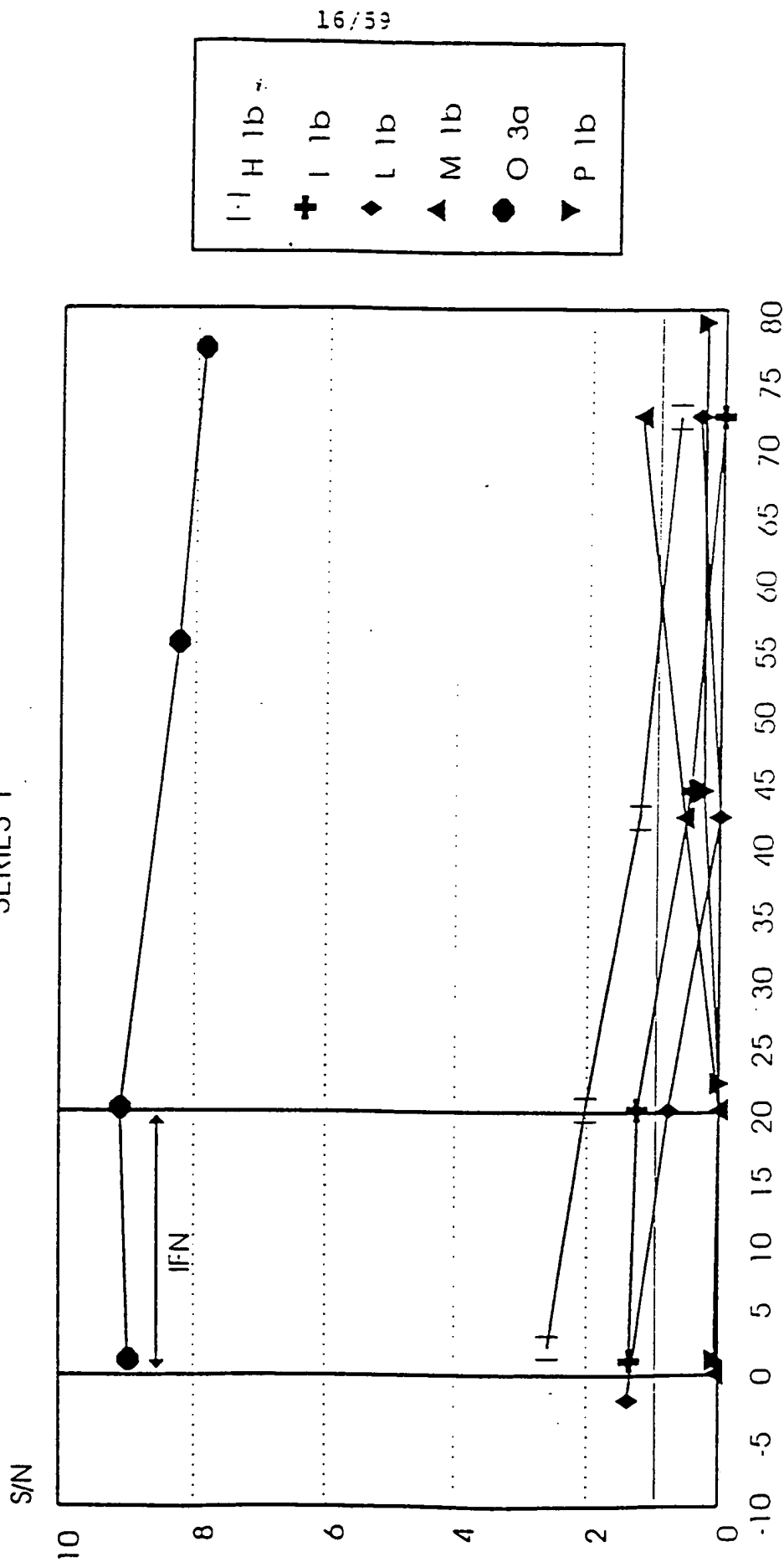


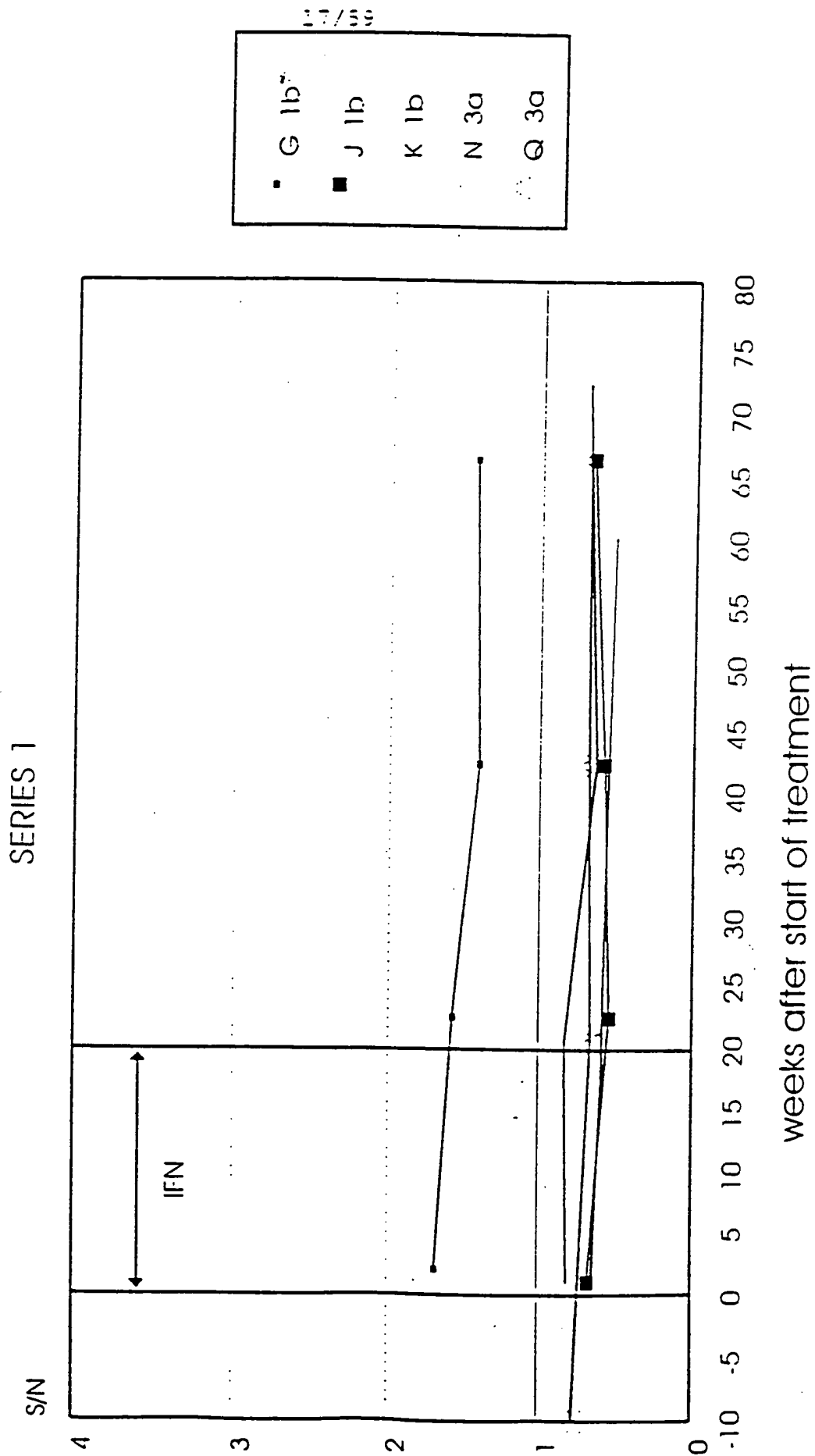
FIGURE 15

Anti-E1 (epitope 1) levels in RESPONDERS to IFN treatment

SERIES 1



nti-E1 (epitope 2) levels in NON-RESPONDERS to IFN treatment



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Anti-E1 (epitope 2) levels in RESPONDERS to IFN treatment

SERIES 1

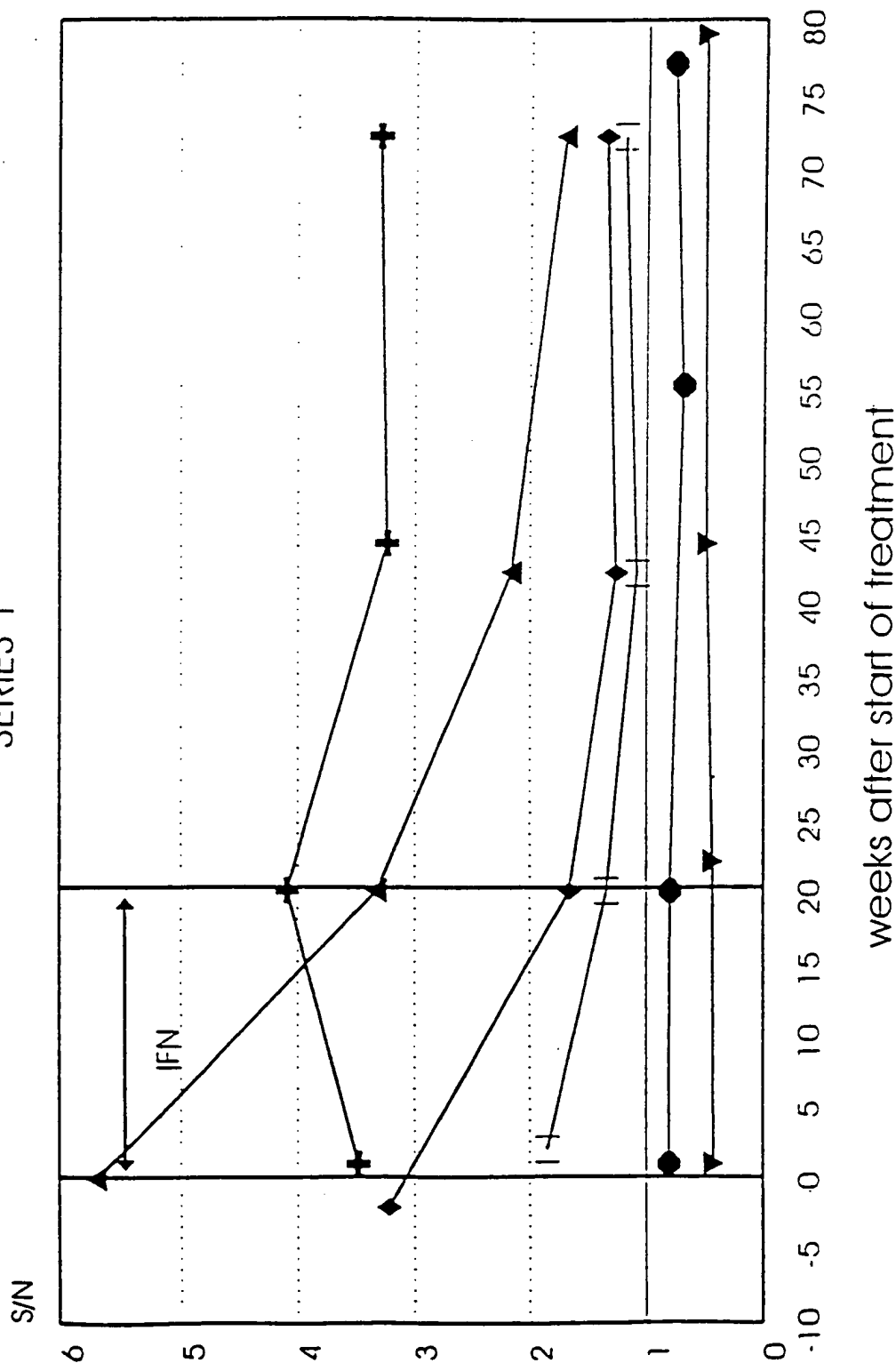
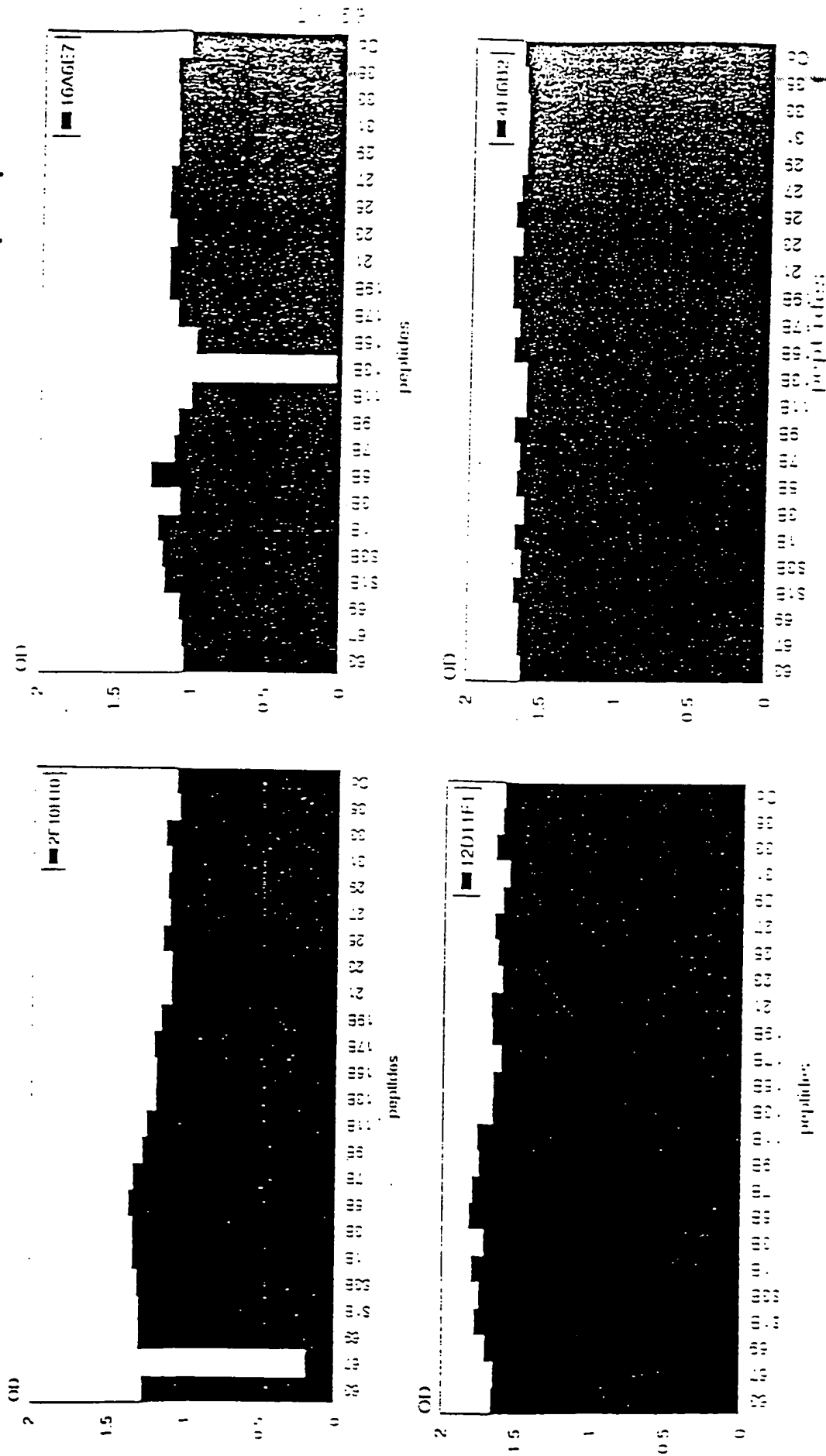


FIGURE 19

Competition of reactivity of anti-E2 Mabs with peptides



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FIGURE 20

Human anti-E2 reactivity competed with peptides

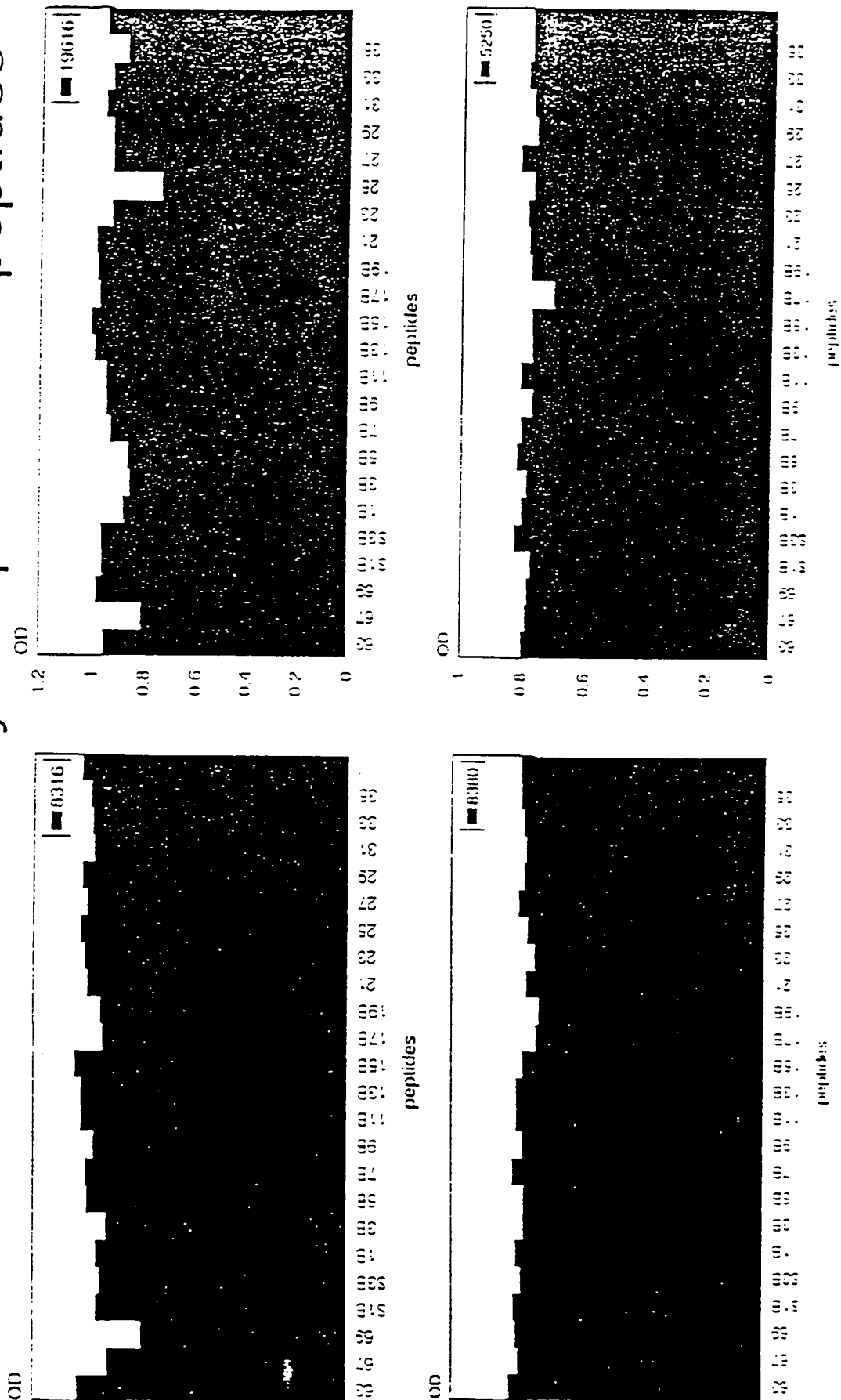


Figure 21

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SEQ ID NO 16 (HCP_r52)

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SEQ ID NO 17 (HCP_r53)

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SEQ ID NO 18 (HCP_r54)

CTATTACCAGTTCATCATCATATCCCA

SEQ ID NO 19 (HCP_r107)

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0969303-070601

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1099999-070601

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09659303-070601

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T0940-074601

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SEQ ID NO 37 (HCCI41)

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TAGGATGTACGTGGGGGGCGTGGAGCACAGGTTTGAAGCCGCATGCAATTGGACTCG
AGGAGAGCGTTGTGACTTGGAGGACAGGGATAGATCAGAGCTTAGCCCGCTGCTGCTG
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SEQ ID NO 41 (HCCI43)

ATGGTGGGGAACTGGGCTAAGGTTTTGGTTGTGATGCTACTCTTTGCCGGCGTCGACG
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GCACTGGGTTCACCAAGACGTGTGGGGGGCCCCCGTGCAACATCGGGGGGGCCGGCA
ACAACACCTTGACCTGCCCCACTGACTGTTTTCGGAAGCACCCCGAGGCCACCTACGC
CAGATGCGGTTCTGGGCCCTGGCTGACACCTAGGTGTATGGTTCATTACCCATATAGG
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096933-070601

SEQ ID NO 43 (HCCI44)

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CAGATGCGGTTCTGGGCCCTGGCTGACACCTAGGTGTATGGTTCATTACCCATATAGG
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SEQ ID NO 45 (HCCL64)

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CCGCGTGTGTCAGGAGGGGCAGCAGCCTCCGATACCAGGGGCCTTGTGTCCCTCTTTAGC
CCCGGGTCTGGCTCAGAAAATCCAGCTCGTAAACACCAACGGCAGTTGGCACATCAAC
AGGACTGCCCTGAACTGCAACGACTCCCTCCAAACAGGGTTCTTTGCCGCACTATTCT
ACAAACACAAATTCAACTCGTCTGGATGCCAGAGCGCTTGGCCAGCTGTGCTCCAT
CGACAAGTTCGCTCAGGGGTGGGGTCCCCTCACTTACACTGAGCCTAACAGCTCGGAC
CAGAGGCCCTACTGCTGGCACTACGCGCCTCGACCGTGTGGTATTGTACCCGCGTCTC
AGGTGTGCGGTCCAGTGTATTGCTTCACCCCGAGCCCTGTTGTGGTGGGGACGACCGA
TCGGTTTGGTGTCCCCACGTATAACTGGGGGGCGAACGACTCGGATGTGCTGATTCTC
AACAACACGCGGCCGCGGAGGCAACTGGTTCGGCTGTACATGGATGAATGGCACT
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T09070-EP95031

SEQ ID NO 47 (HCCI65)

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GGTTCTGGAGGACGGCGTGAACTATGCAACAGGGAATTTGCCCGGTTGCTCTTTCTCT
ATCTTCCTCTTGGCTTTGCTGTCTGTCTGACCGTTCCAGCTTCGCTTATGAAGTGCG
CAACGTGTCCGGGATGTACCATGTACGAACGACTGCTCCAACCTCAAGCATTGTGTAT
GAGGCAGCGGACATGATCATGCACACCCCCGGGTGCGTGCCCTGCGTTCGGGAGAAC
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GGGCCCCATTGGGGAGTCCTGGCGGGCCTCGCCTACTATTCCATGGTGGGGAACTGGGC
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ACGTGCAATACCTGTACGGTGTAGGGTCGGCGGTTGTCTCCCTTGTCAATCAAATGGGA
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TGATGCTGCTGATAGCTCAAGCTGAGGCCGCTTAGAGAACCTGGTGGTCCCTCAATGC
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SEQ ID NO 49 (HCC166)

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CTCAGCCCCGGGTACCCTTGGCCCCCTCTATGGCAATGAGGGCATGGGGTGGGCAGGATG
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AGCGTCCCCACCACGACAATACGACGCCACGTCGATTTGCTCGTTGGGGCGGCTGCTT
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GCGGGGGGCCATTGGGGAGTCCCTGGCGGGCCCTCGCCTACTATTCCATGGTGGGGAAC
GGGCTAAGGTTTTGGTTGTGATGCTACTCTTTGCCGGCGTCGACGGGCATACCCGCGT
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09869403.070601

TCGGCTCAGAAAATCCAGCTCGTAAACACCAACGGCAGTTGGCACATCAACAGGACT
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GCTGCCTGGTACATCAAGGGCAGGCTGGTCCCTGGTGCGGCATACGCCTTCTATGGCG
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0969303-070601

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Figure 22

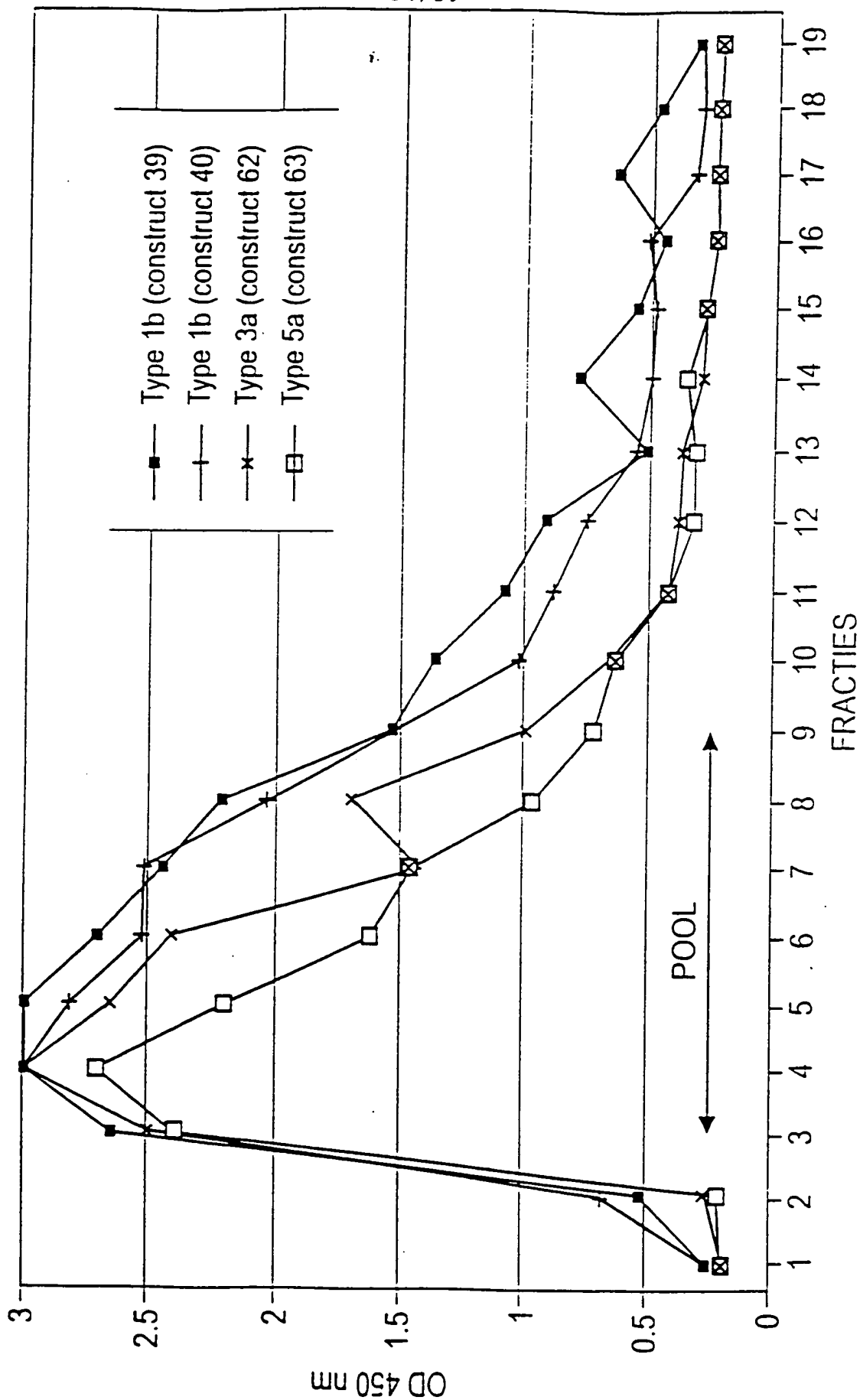
OD measured at 450 nm
construct

Fraction	volume dilution	39 Type 1b	40 Type 1b	62 Type 3a	63 Type 3a
START	23 ml 1/20	2.517	1.954	1.426	1.142
FLOW THROUGH	23 ml 1/20	0.087	0.085	0.176	0.120
1	0.4 ml 1/200	0.102	0.051	0.048	0.050
2		0.396	0.550	0.090	0.067
3		2.627	2.603	2.481	2.372
4		3	2.967	3	2.694
5		3	2.810	2.640	2.154
6		2.694	2.499	1.359	1.561
7		2.403	2.481	0.347	1.390
8		2.176	1.970	1.624	0.865
9		1.461	1.422	0.887	0.604
10		1.286	0.926	0.543	0.519
11		0.981	0.781	0.294	0.294
12		0.812	0.650	0.249	0.199
13		0.373	0.432	0.239	0.209
14		0.653	0.371	0.145	0.184
15		0.441	0.348	0.151	0.151
16		0.321	0.374	0.098	0.106
17		0.525	0.186	0.099	0.108
18		0.351	0.171	0.083	0.090
19		0.192	0.164	0.084	0.087

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FIG. 10



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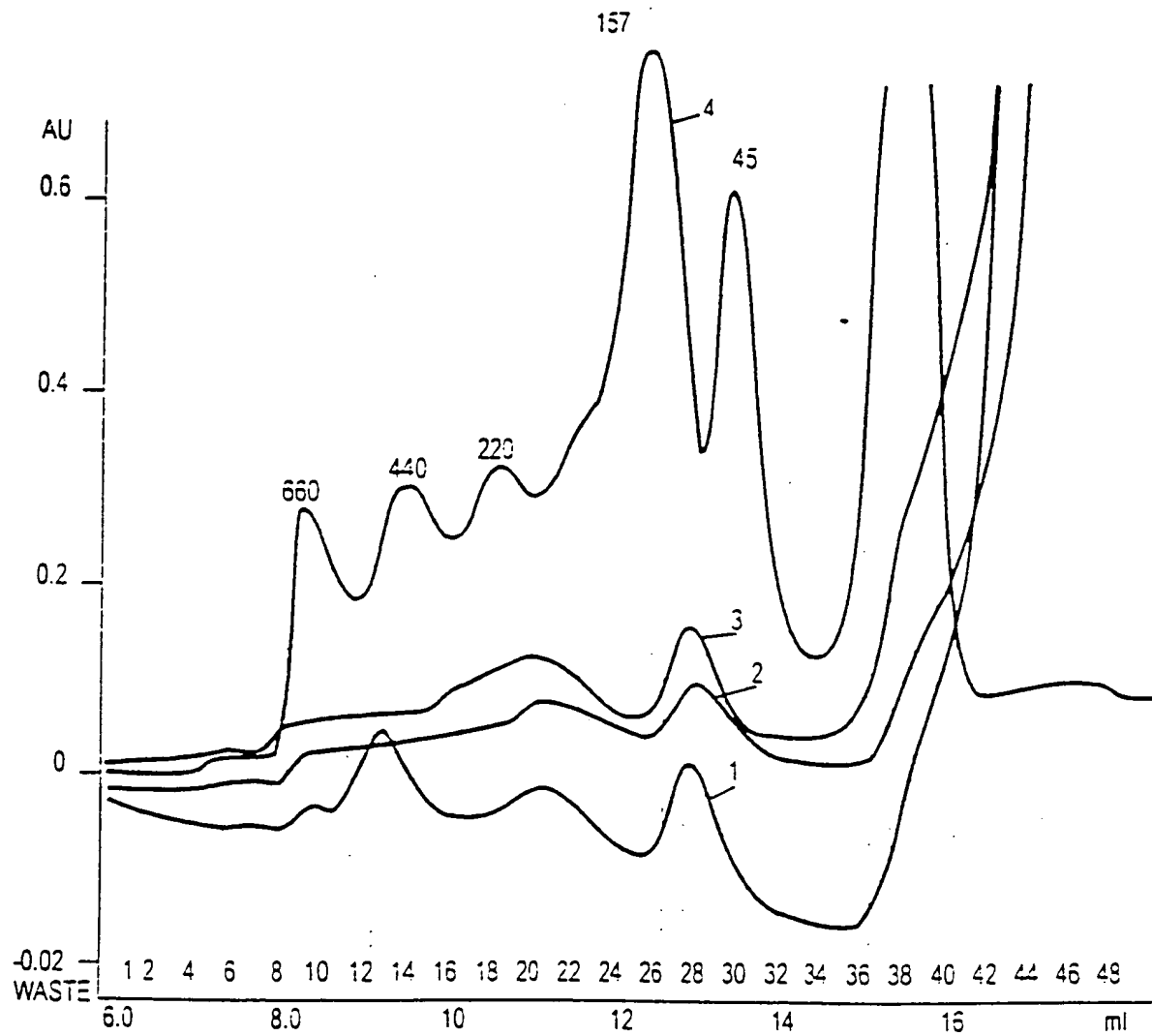


FIGURE 25

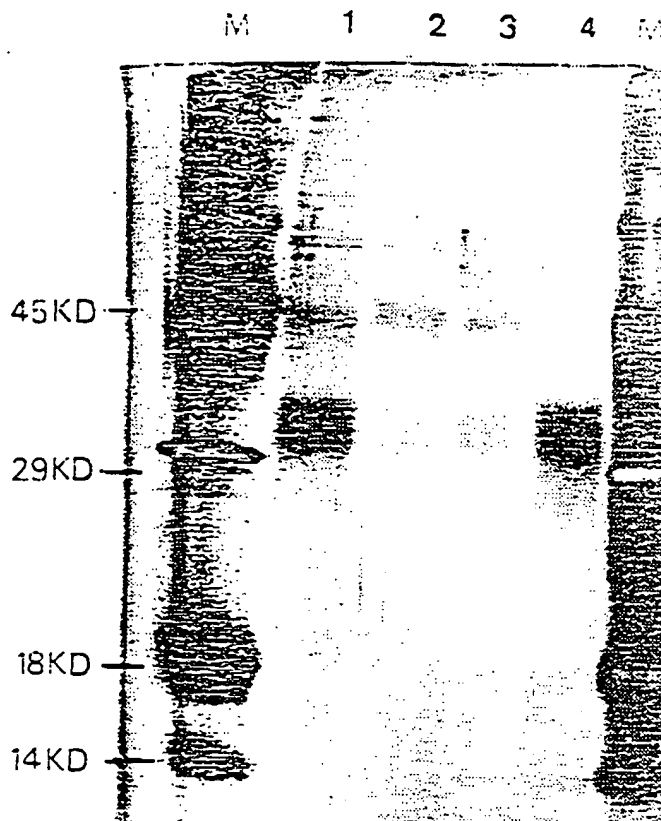


Figure 26

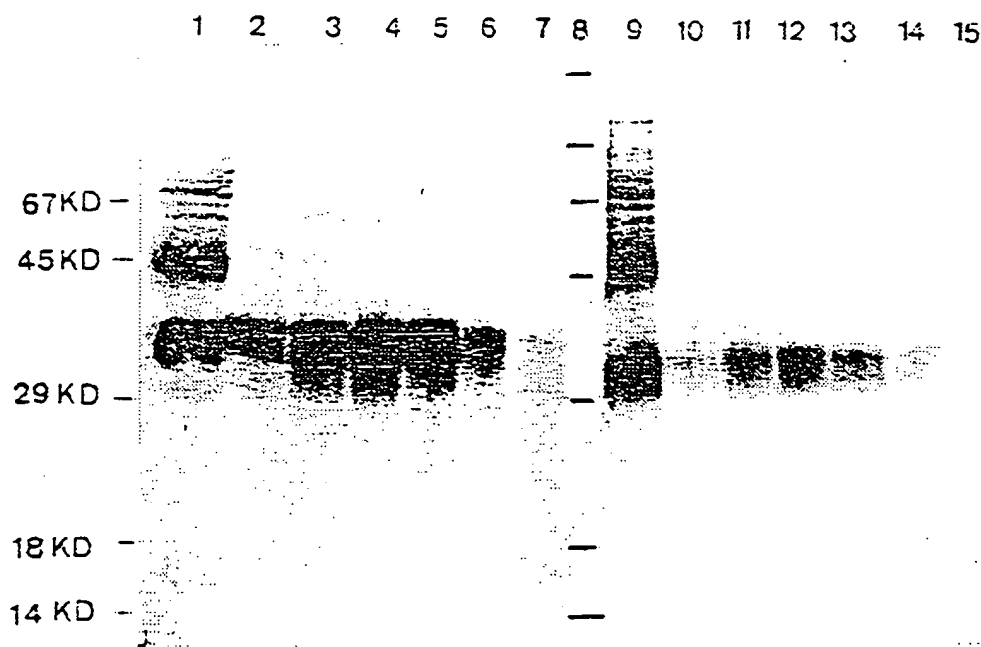


Figure 27

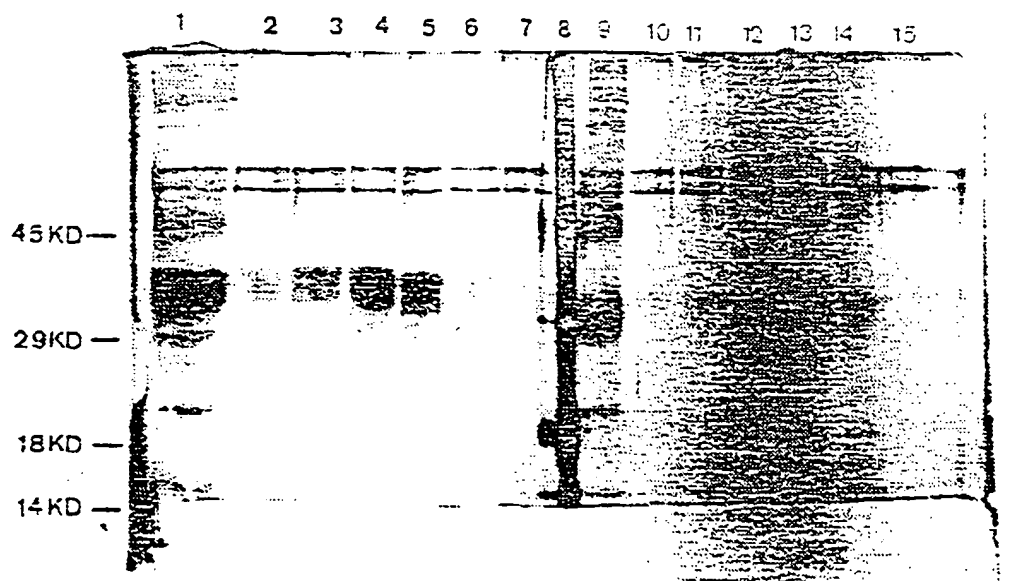
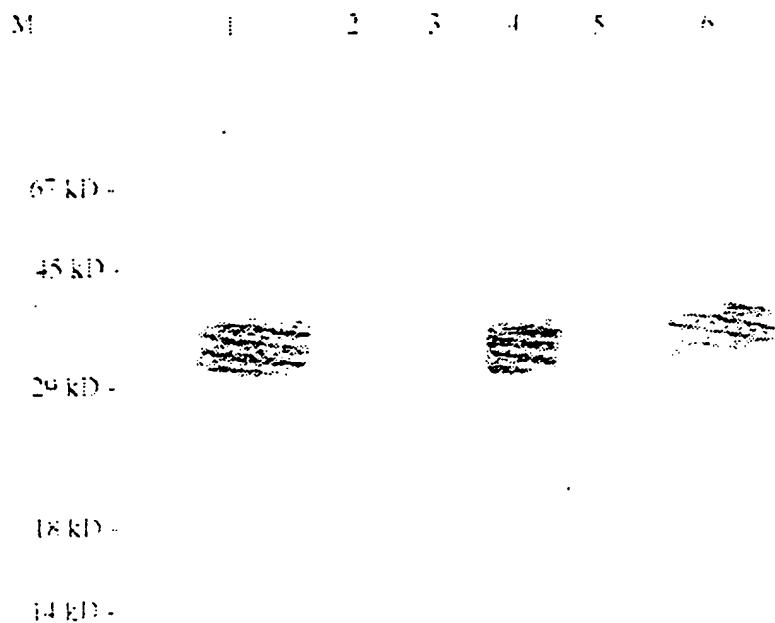


Figure 2E



Lane 1: Crude Lysate
Lane 2: Flow through Lentil Chromatography
Lane 3: Wash with EMPIGEN Lentil Chromatography
Lane 4: Eluate Lentil Chromatography
Lane 5: Flow through during concentration lentil eluate
Lane 6: Pool of E1 after Size Exclusion Chromatography

Figure 29: Western Blot Analysis with anti-E1 mouse monoclonal 5E1A10

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109040" E0266860

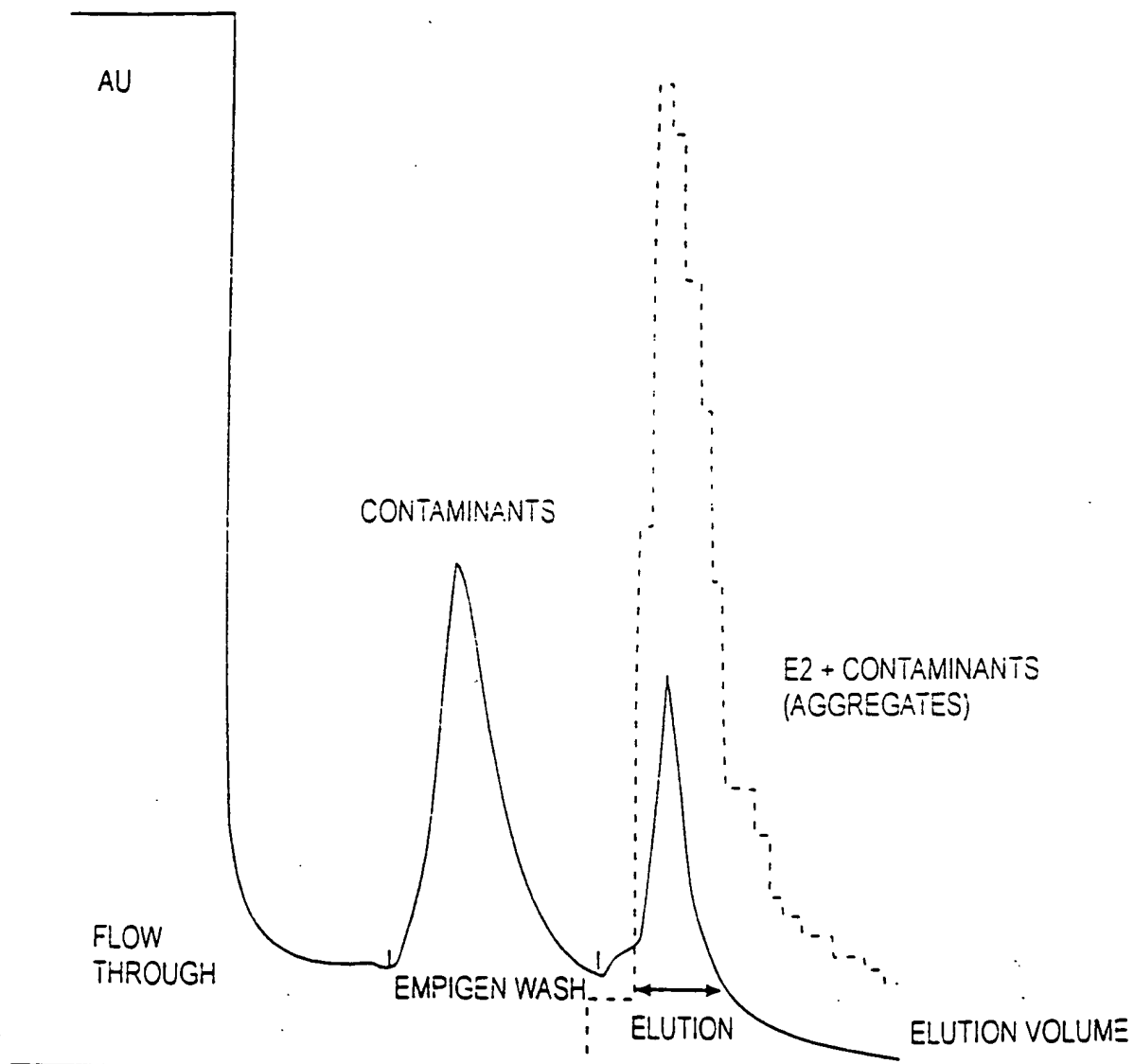
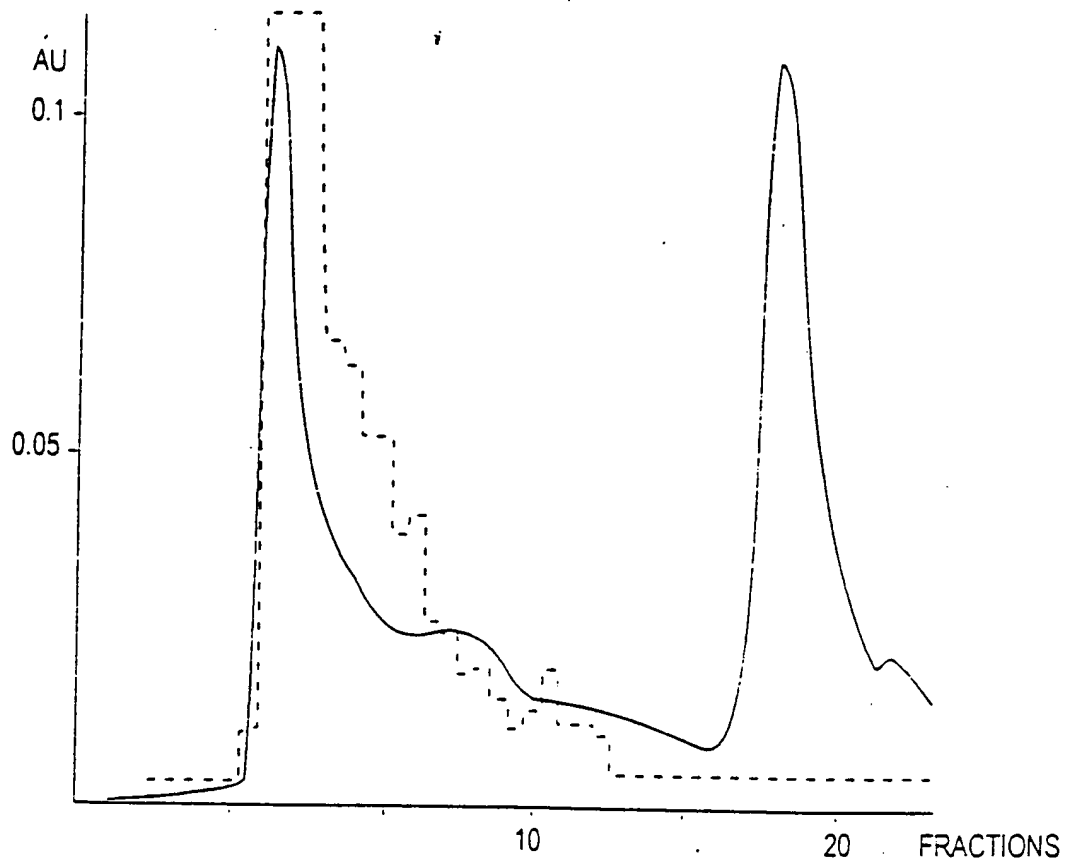


FIGURE 30

A: NON - REDUCED

E2 - CONTAMINANTS (AGGREGATES)



B: REDUCED

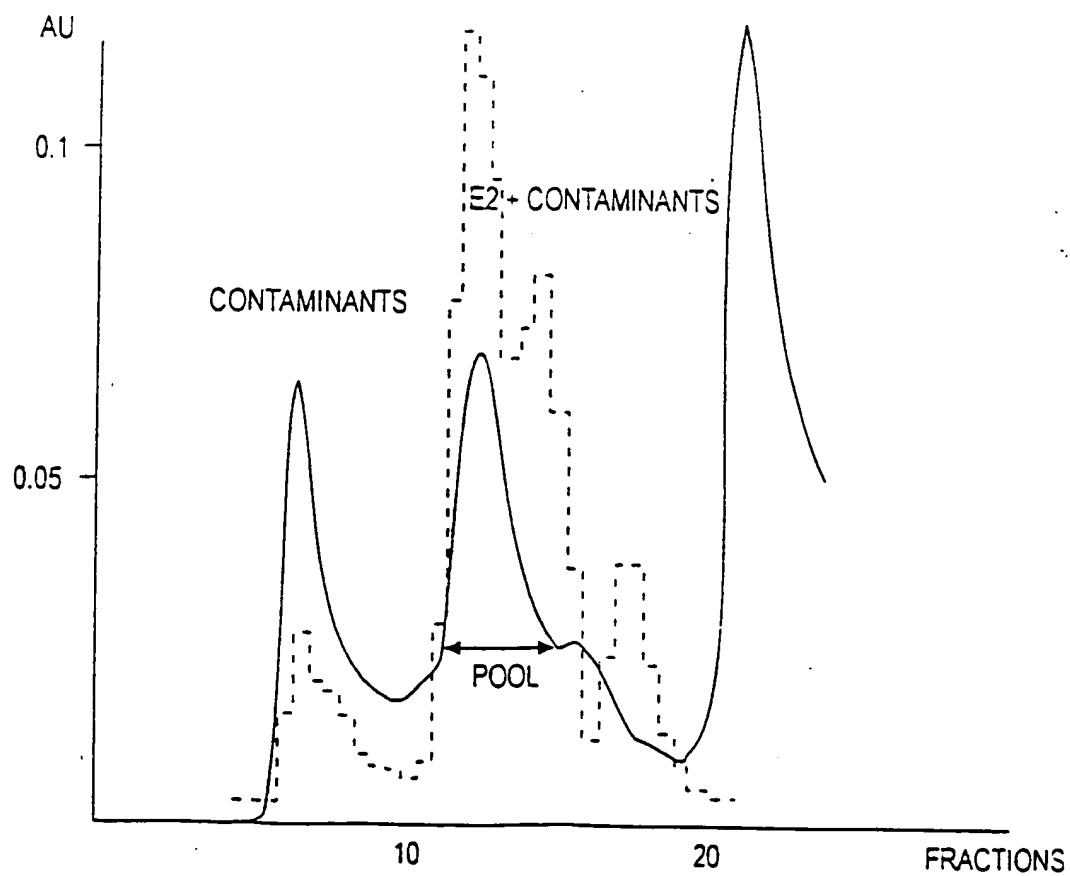


FIGURE 31

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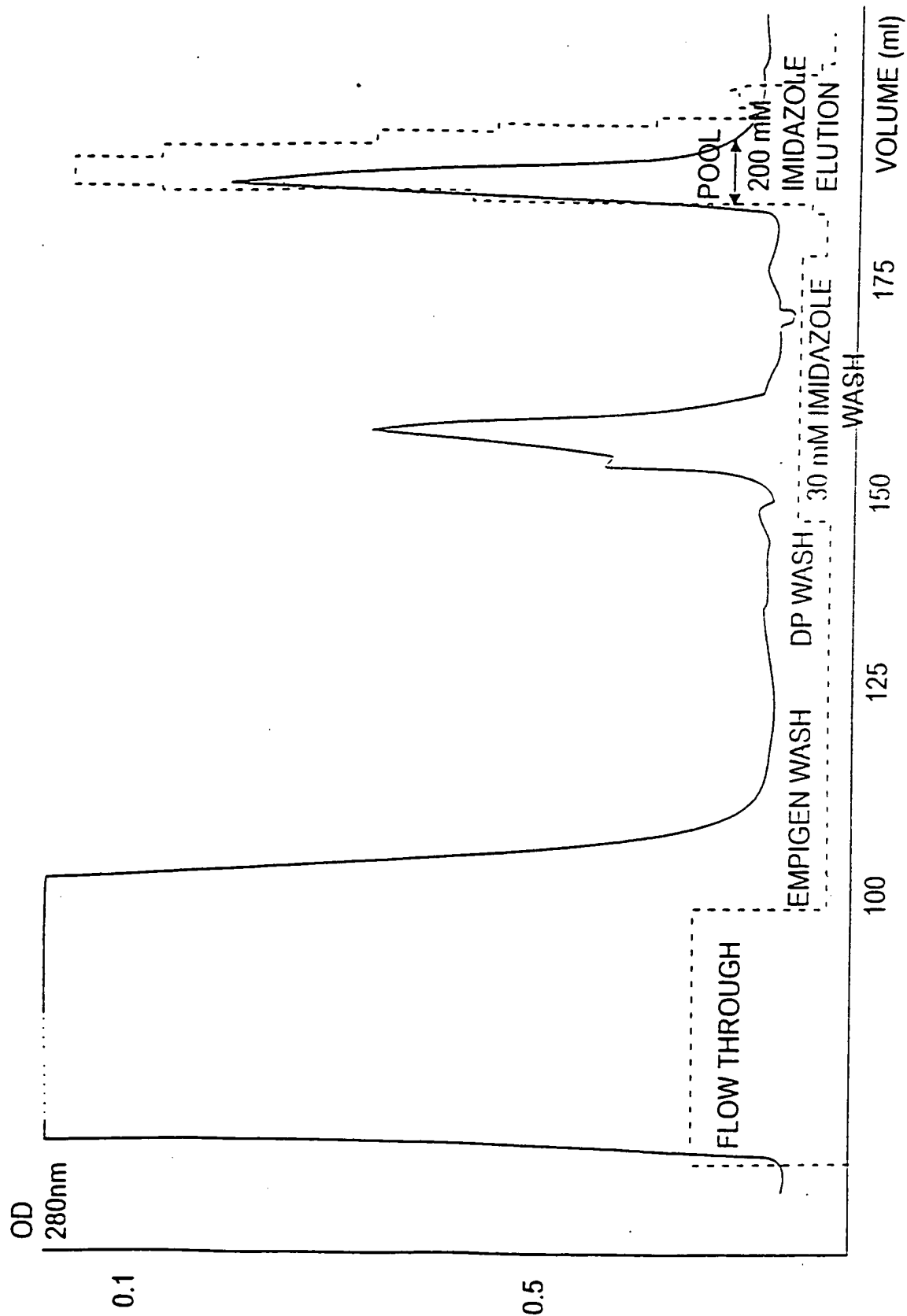
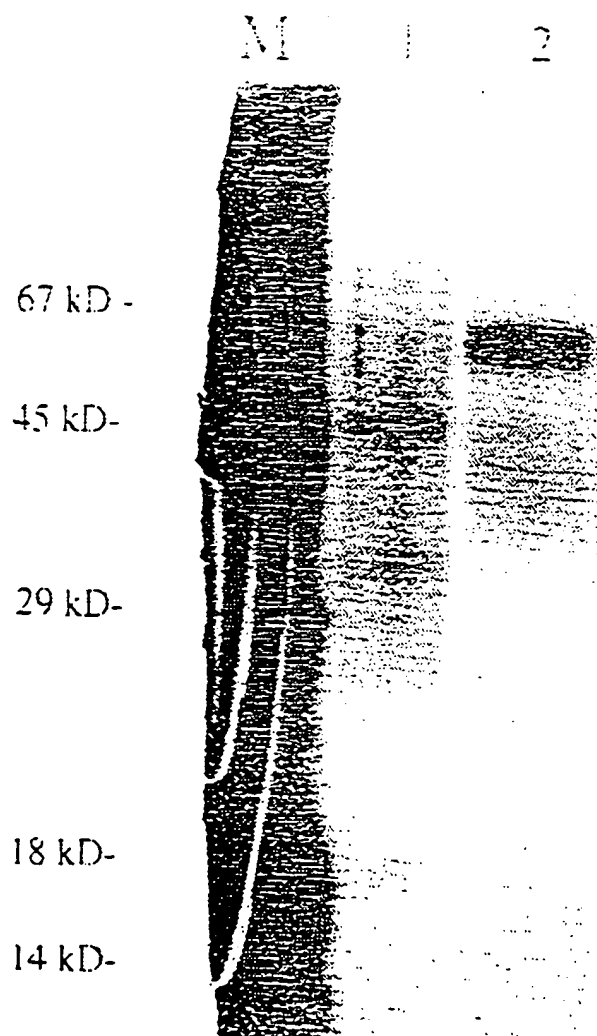


FIGURE 32

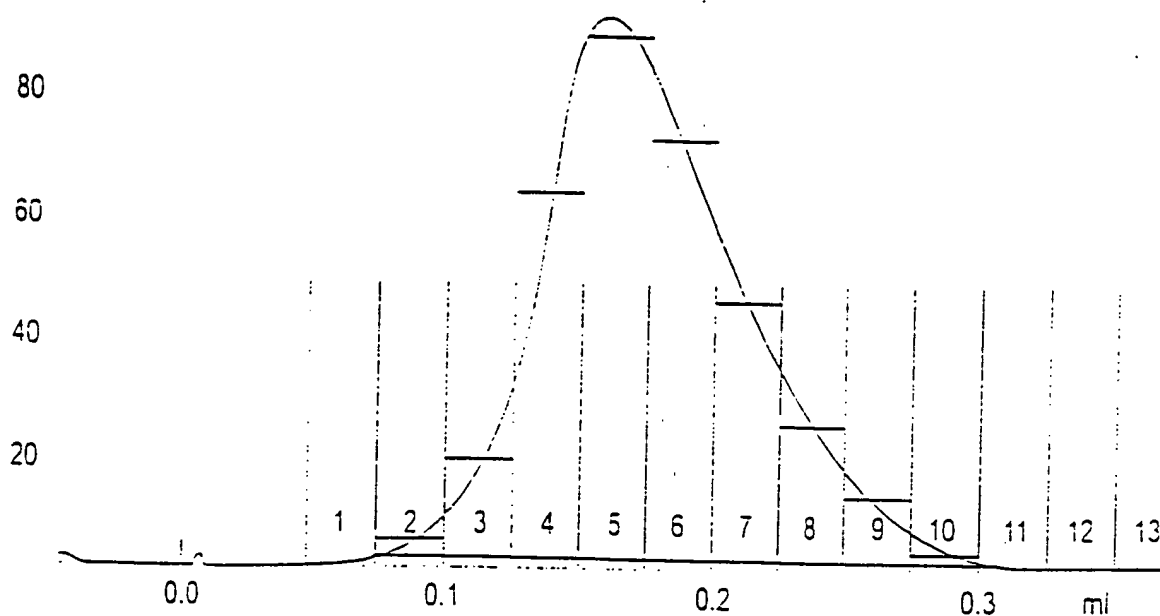
109040" E066660

FIGURE 33:
SILVER STAIN OF PURIFIED E2



1. 30 mM IMIDAZOLE WASH Ni-IMAC
2. 0.5 μ g E2

45 59 Figure 34



No.	Ret. (ml)	Peak start (ml)	Peak end (ml)	Dur (ml)	Area (ml*mAU)	Height (mAU)
1	-0.45	-0.46	-0.43	0.04	0.0976	4.579
2	1.55	0.75	3.26	2.51	796.4167	889.377
3	3.27	3.26	3.31	0.05	0.0067	0.224
4	3.35	3.32	3.33	0.02	0.0002	0.018

Total number of detected peaks = 4

Total Area above baseline = 0.796522 ml*mAU

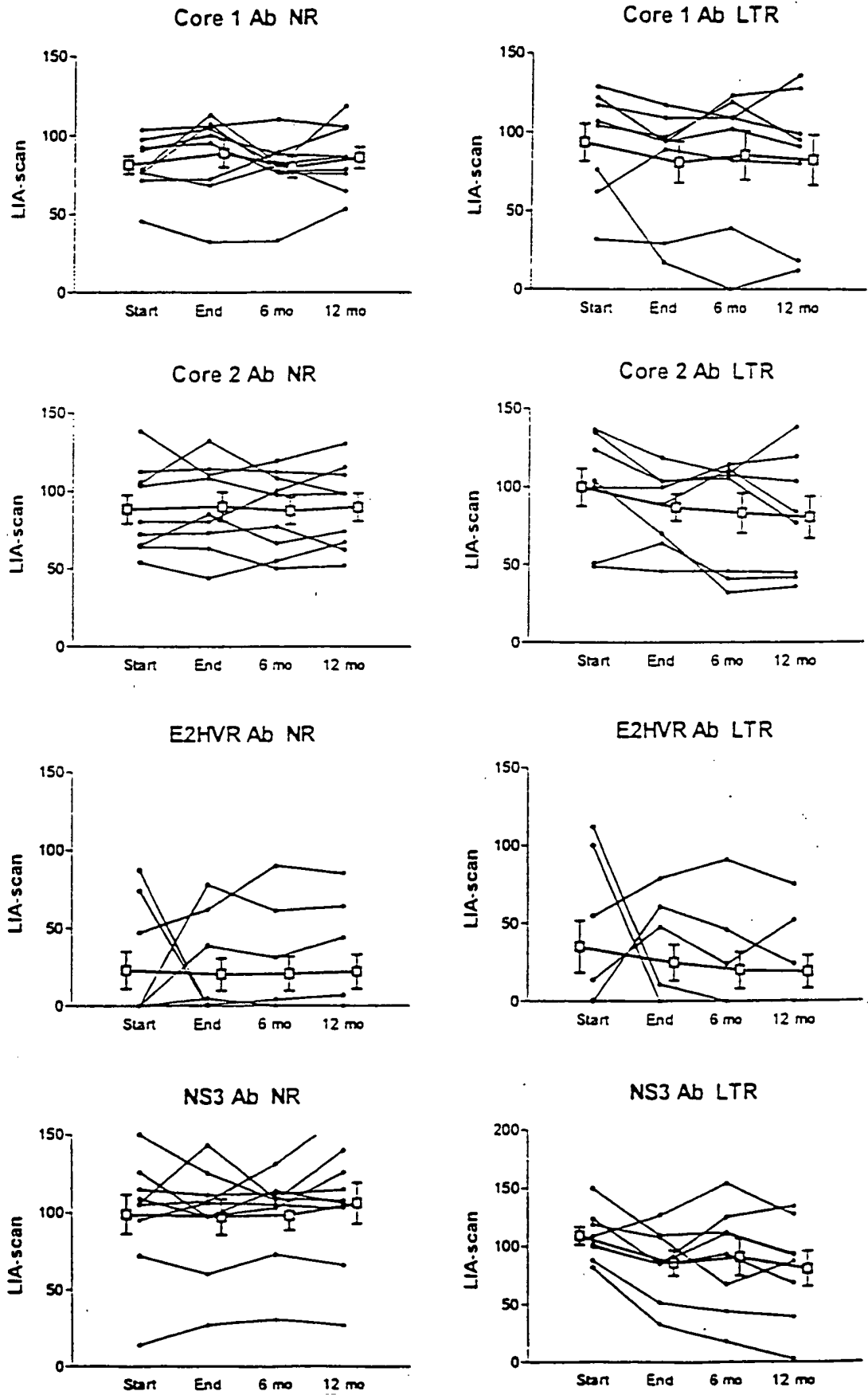
Total area in evaluated peaks = 0.796521 ml*mAU

Ratio peak area / total area = 0.999999

Total peak duration = 2.613583 ml

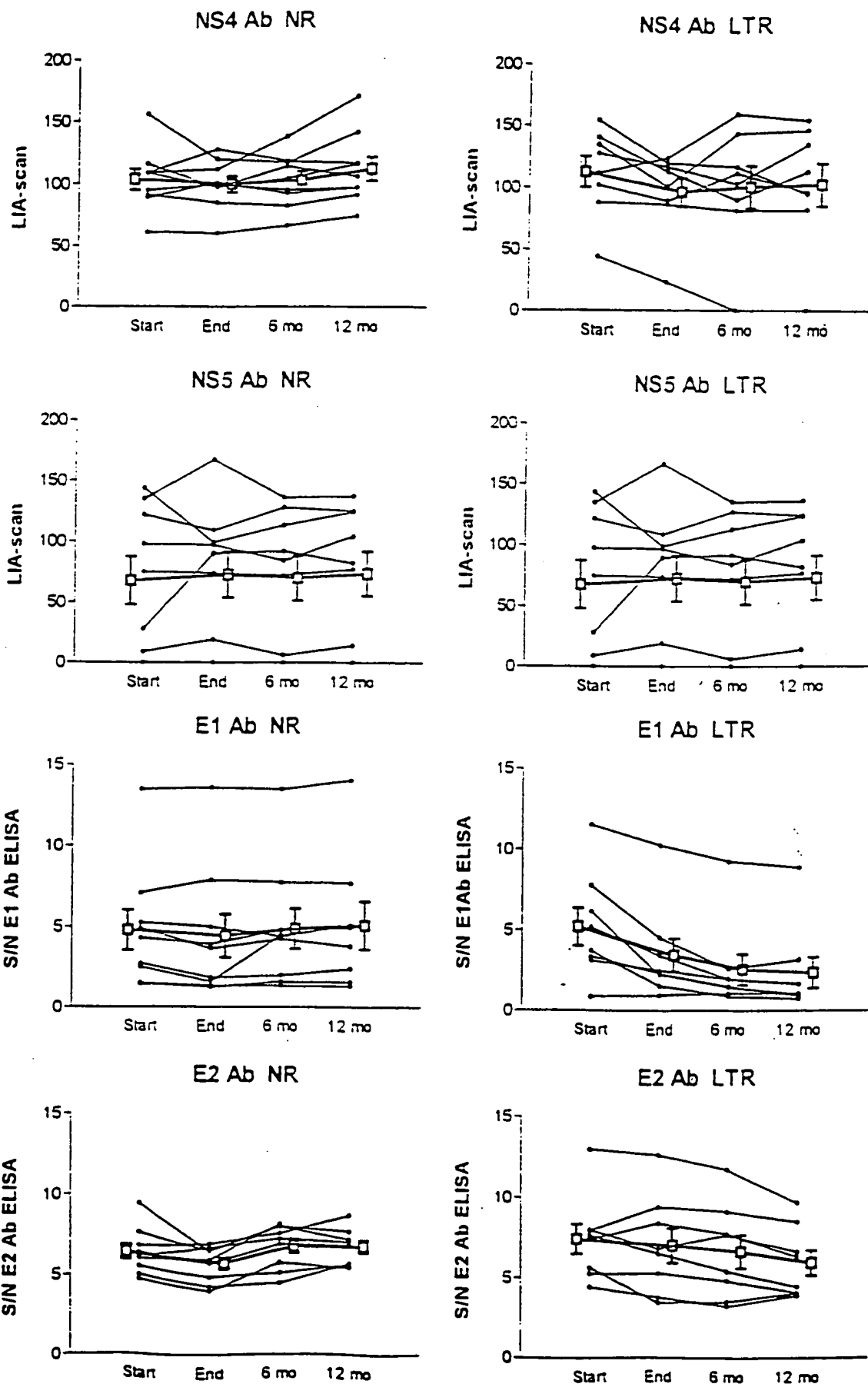
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FIGURE 35A



T09070" E0F66860

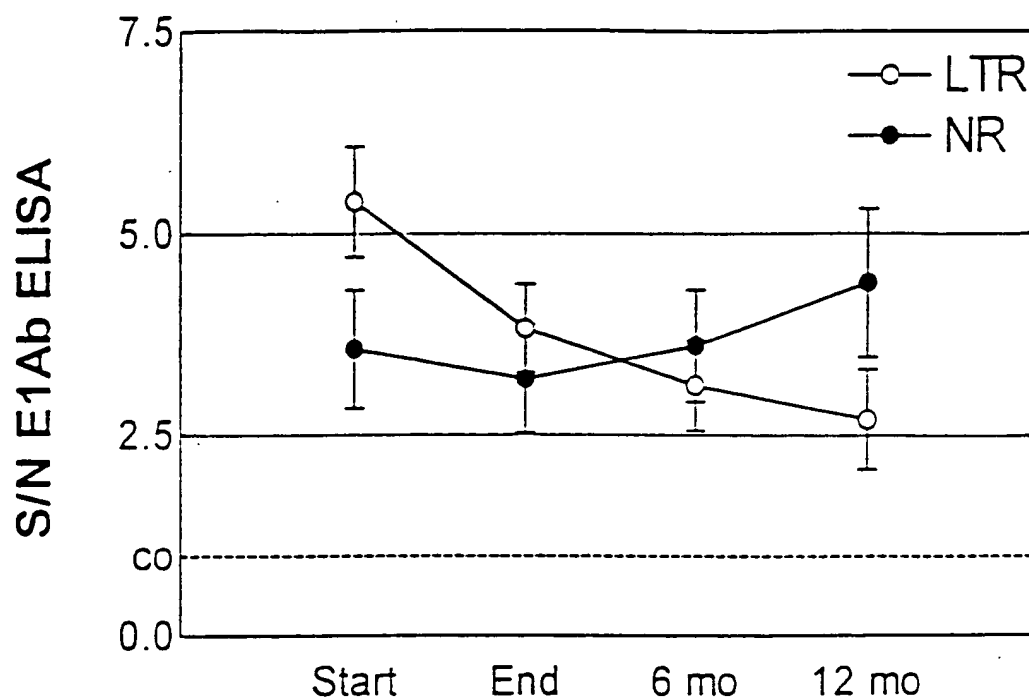
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FIGURE 35B

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Figure 36

E1 Ab



E2 Ab

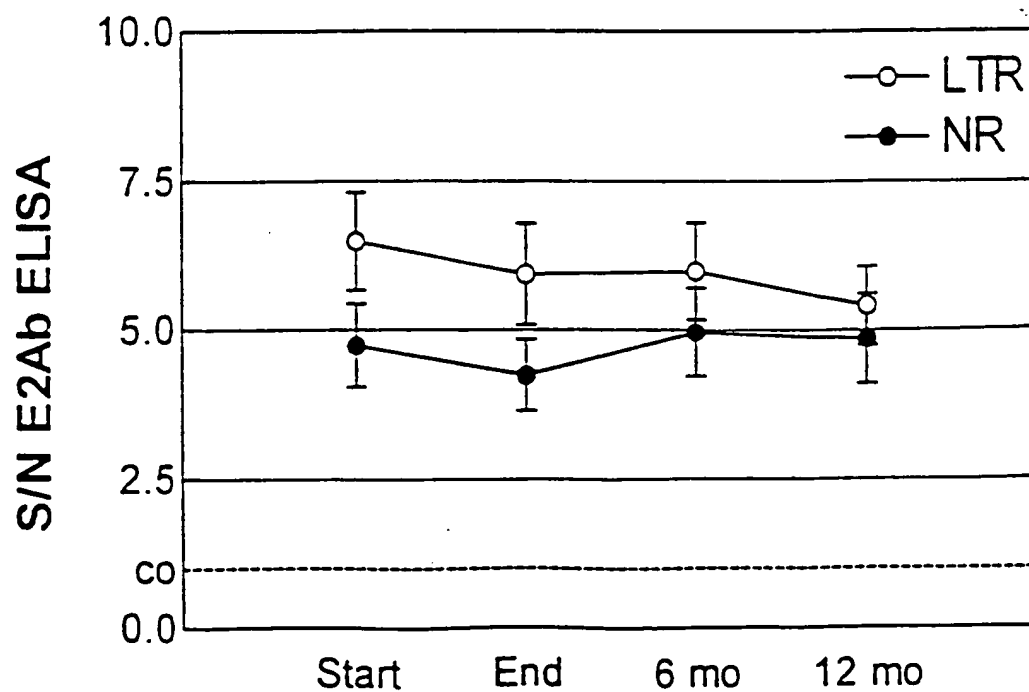


FIGURE 37

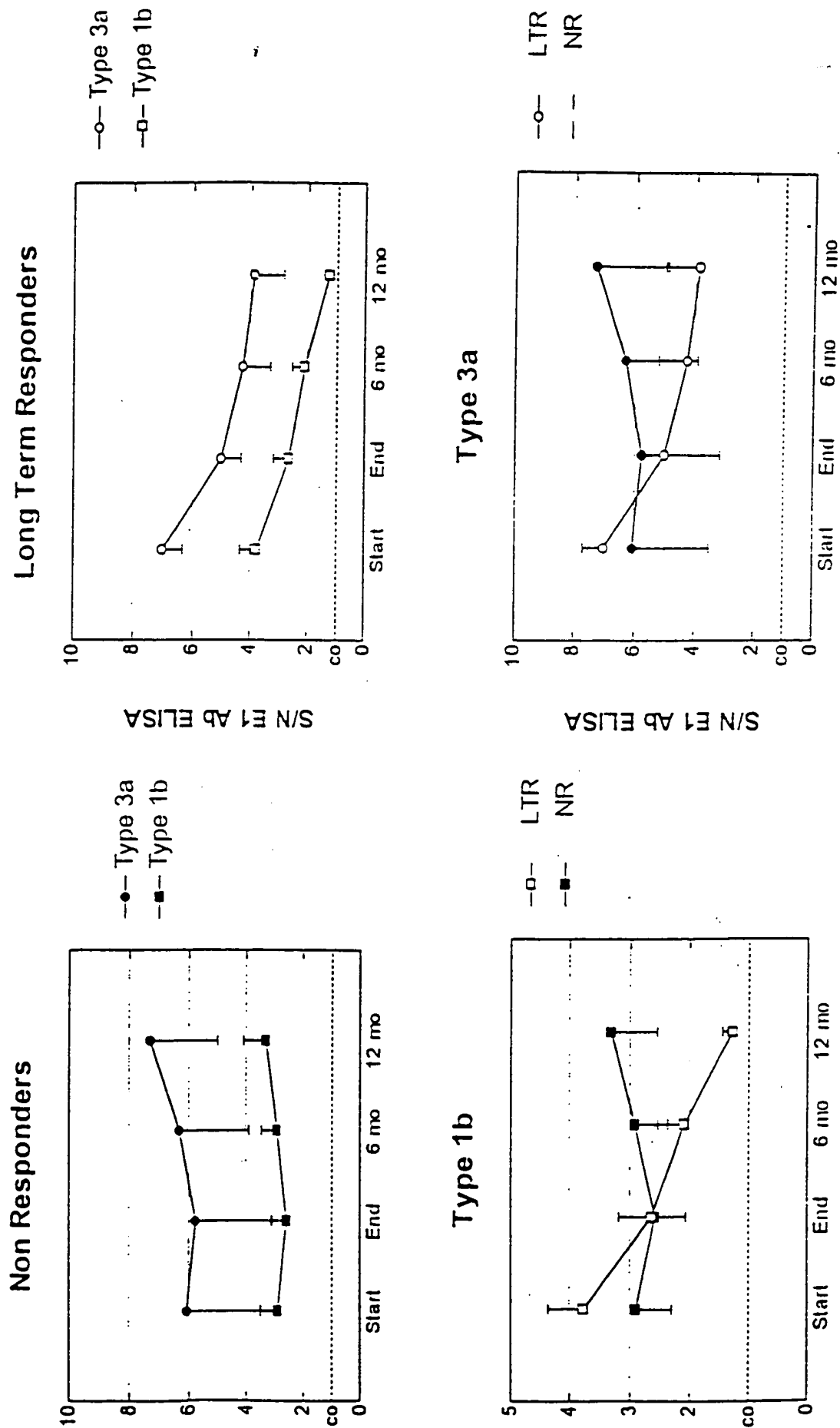
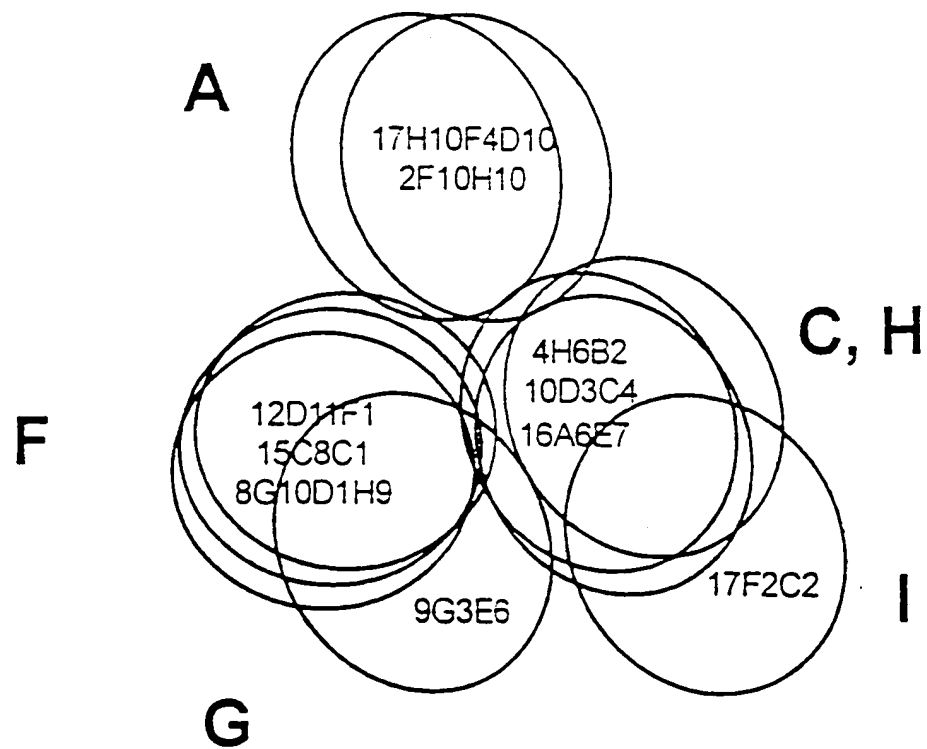


Figure 38

Relative Map Positions of
anti-E2 monoclonal antibodies



0989303-070501

109070" E0266860

PARTIAL DEGLYCOSYLATION OF HCV E1 ENVELOPE PROTEIN

Endoglycosidase H Glycopeptidase F
(Endo H) (PNGase F)

0µg 0.6µg 6µg 60µg 0.6µg 6µg 0.04µg 0.4µg 4µg 40µg 400µg

106.0
80.0
49.5
32.5
27.5
18.5

↑
↑
↑
↑
↑
↑
↑

Figure 39

PARTIAL TREATMENT OF HCV E2/E2s ENVELOPE PROTEINS BY PNGase F

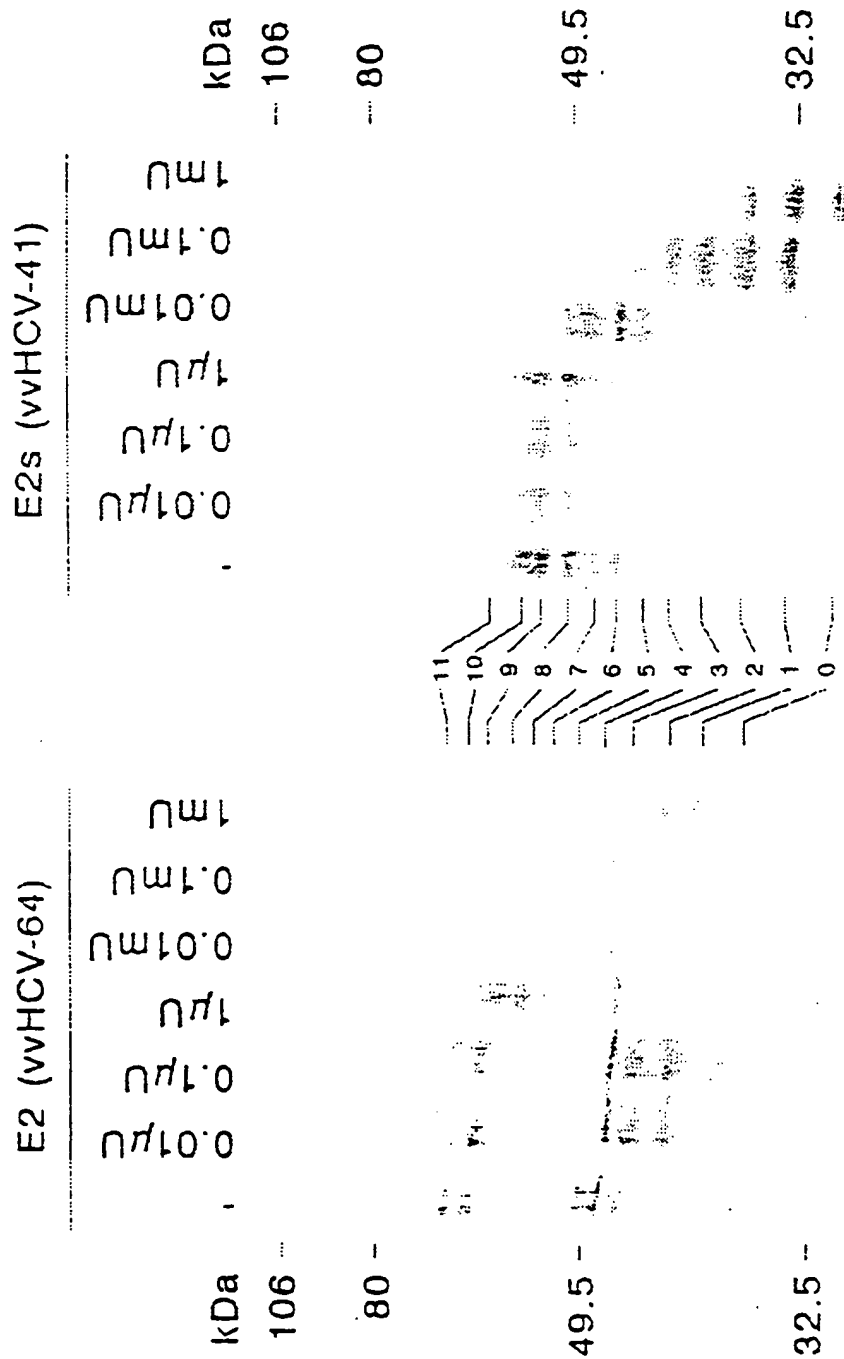


Figure 40

Fig. 41 *In Vitro* Mutagenesis of IICV E1 glycoprotein

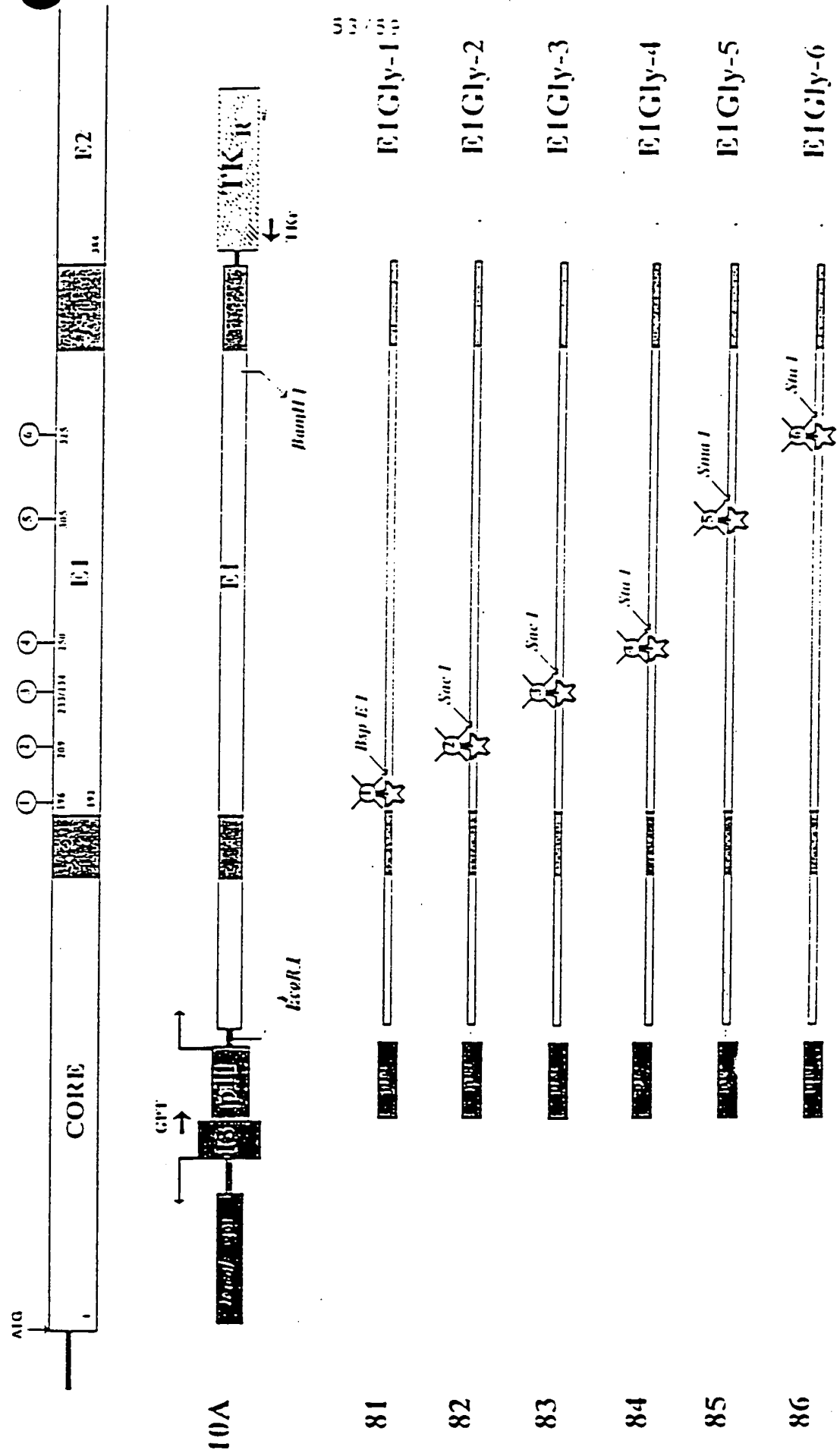
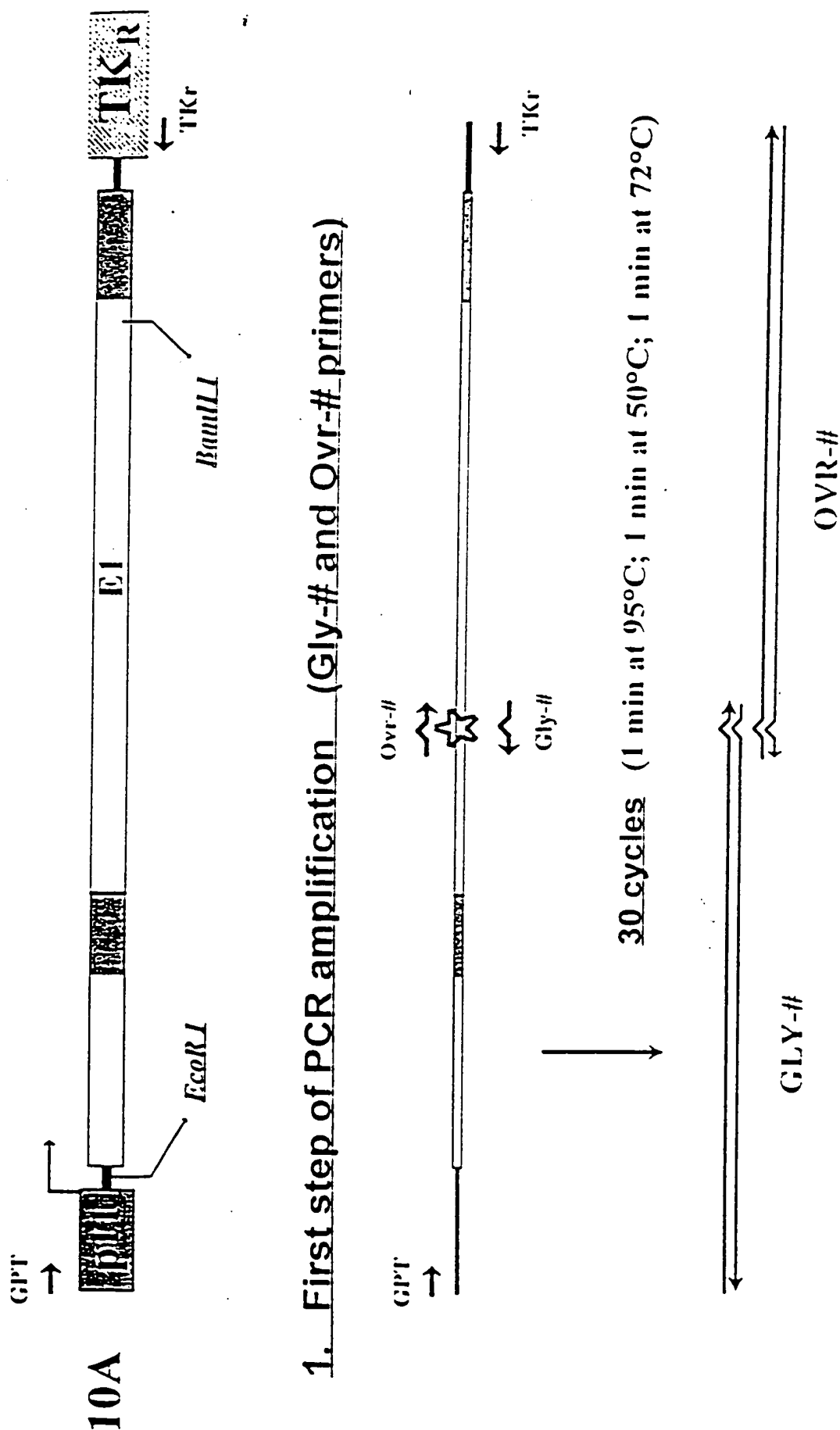
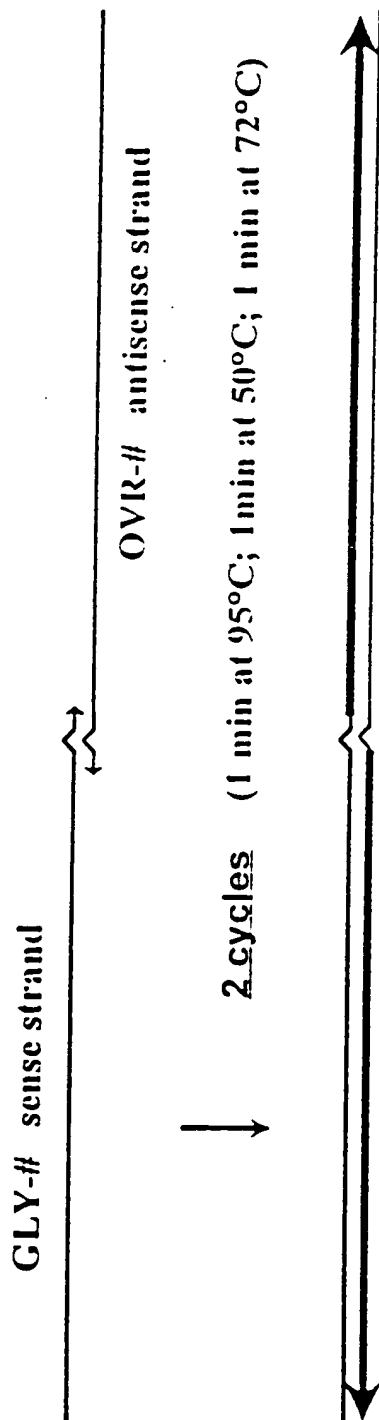


Fig. 42A In Vitro Mutagenesis of HCV E1 glycoprotein



2. Overlap extension and nested PCR Fig. 42B

a. Overlap extension



b. Nested PCR amplification (GPT-2 and TKR-2 primers)

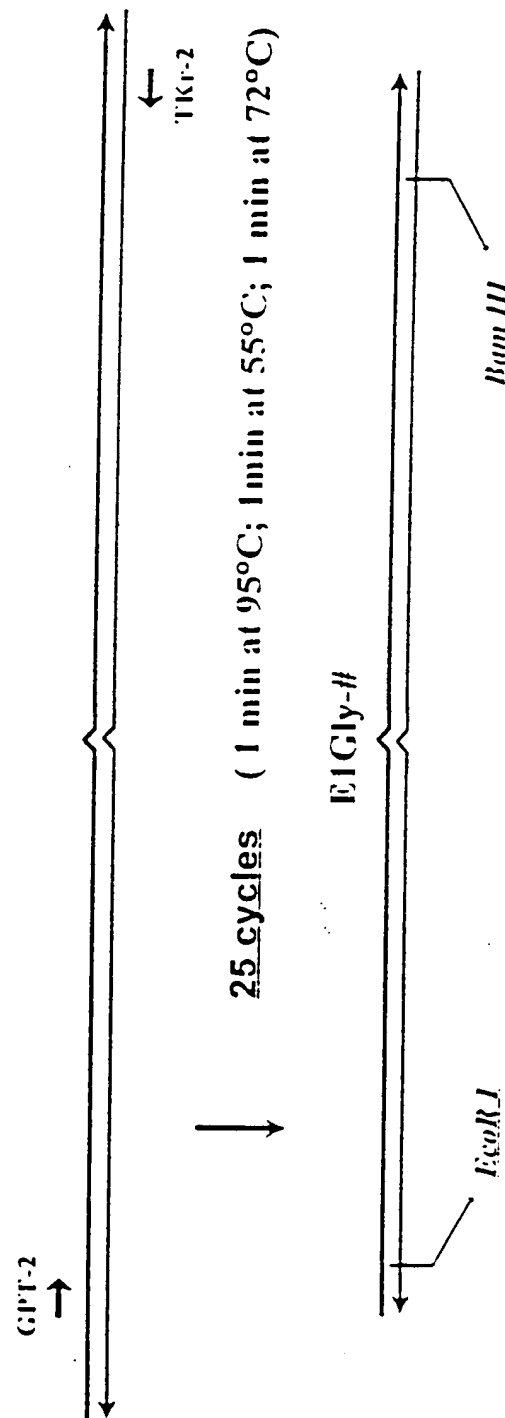
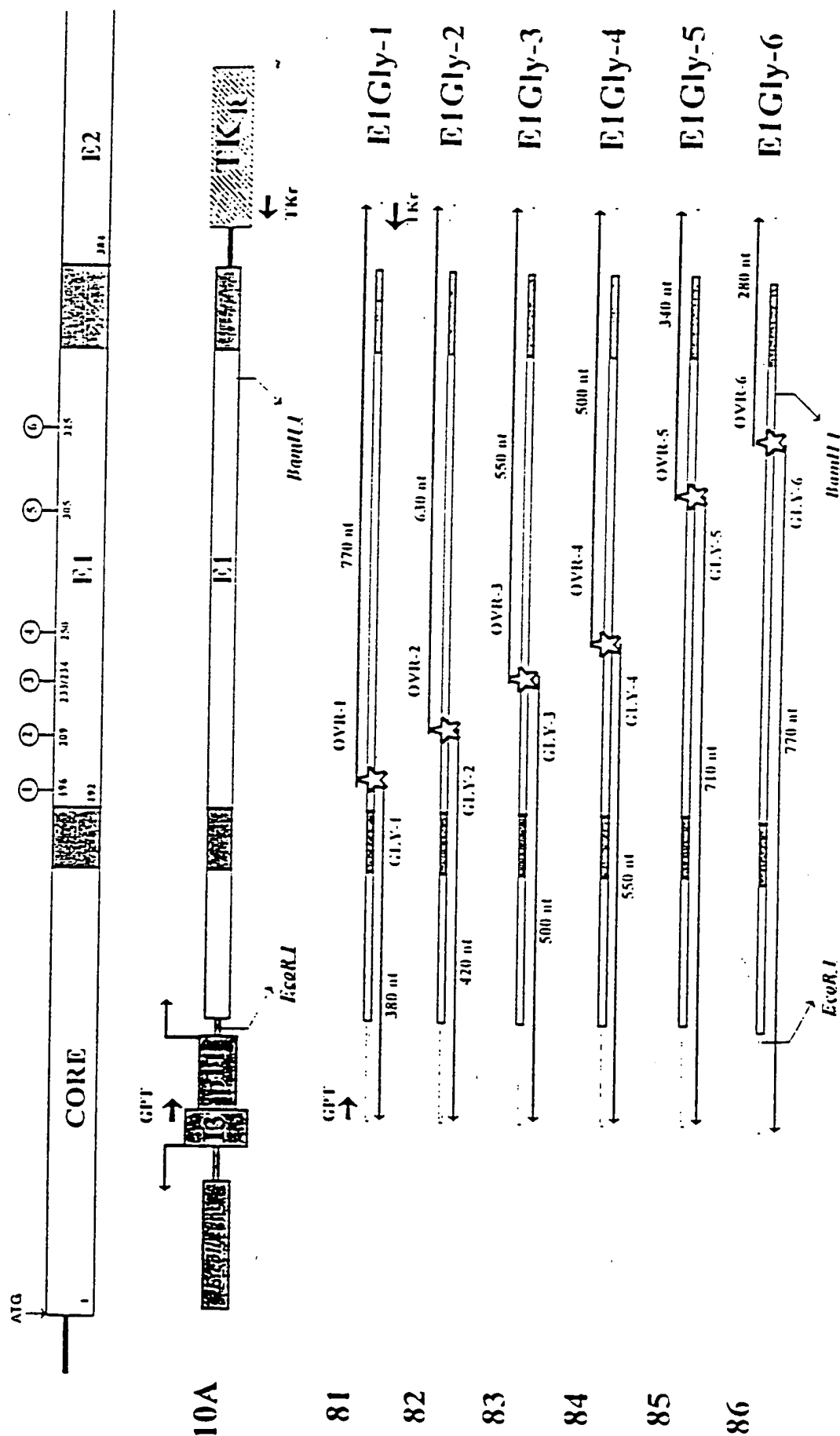


Fig. 43 *In Vitro* Mutagenesis of HCV E1 glycoprotein



HeLa cells										RK 13 cells												
<div>1234567</div>										<div>21345678</div>												
80.0	—									—	80.0										—	80.0
49.5	—									—	49.5										—	49.5
32.5	—									—	32.5										—	32.5
27.5	—									—	27.5										—	27.5
18.5	—									—	18.5										—	18.5

Figure 44A

098666860

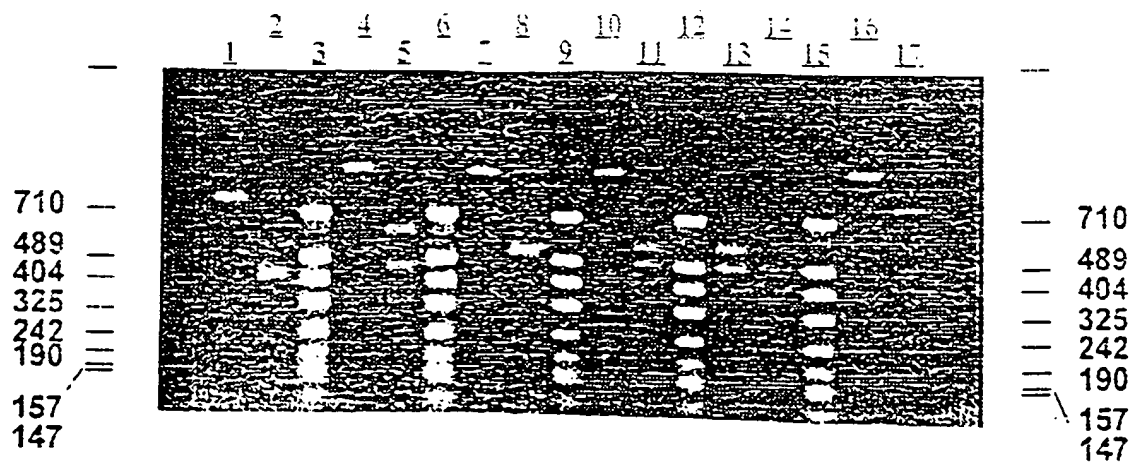


Figure 4-18

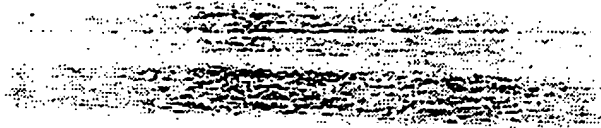


Figure 45

kDa
 — 119
 — 67
 — 43
 — 29
 — 18



Figure 46

109070- E0E66860